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December 28, 2005, 13:43:04; Search time 147.538 Seconds (without alignments) 375.236 Million cell updates/sec
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668
1 QVQLVBSGGGVVQPGRSLRL.....NYYYYGMDVWGQGTTVTVSS 126
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Listing first 45 summaries
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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25 577.5 86.5 1125 8 ADP22316 27 577 86.4 115 5 AAM51164 28 576 86.2 118 5 AAM51167 28 575.5 86.2 118 5 AAM51167 30 575.5 86.2 118 5 AAM51167 31 573.5 86.2 118 6 ADP22300 32 572.5 85.7 25.2 5 ABP5720 33 572.5 85.7 25.2 5 ABP5720 34 572.5 85.7 25.2 5 ABP5720 35 572.5 85.7 25.2 5 ABP5720 36 572.5 85.7 25.2 5 ABP5721 37 572.5 85.7 25.2 5 ABP57320 38 571.5 85.6 122 8 ADP22128 38 571.5 85.6 123 8 ADR88117 41 570.5 85.4 122 7 ADE28451 42 570.5 85.4 125 7 ADE28451 43 570.5 85.4 125 7 ADE28475 44 570.5 85.4 125 7 ADE28475 45 569.5 85.4 127 9 AEB7751 47 570.5 85.4 127 9 AEB7751 48 570.5 85.4 127 9 AEB7751 49 570.5 85.4 127 9 AEB7751 40 570.5 85.4 127 9 AEB7751 41 570.5 85.4 127 9 AEB7751 42 570.5 85.4 470 7 ADE28475 43 570.5 85.4 127 9 AEB7751 44 570.5 85.4 127 9 AEB7751 45 569.5 85.3 122 9 AEB7751 46 570.5 85.4 127 9 AEB7751 47 570.5 85.4 127 9 AEB7751 48 570.5 85.4 127 9 AEB7751 49 570.5 85.4 127 9 AEB7751 40 570.5 85.4 127 9 AEB7751 41 570.5 85.4 127 9 AEB7751 42 570.5 85.4 127 9 AEB7751 43 570.5 85.4 127 9 AEB7751 44 570.5 85.4 127 9 AEB7751 45 569.5 85.3 122 9 AEB7751 46 570.5 85.4 127 9 AEB7751 47 570.5 85.4 127 9 AEB7751 48 570.5 85.4 127 9 AEB7751 49 570.5 85.4 127 9 AEB7751 40 570.5 85.4 127 9 AEB7751 41 570.5 85.4 127 9 AEB7751 42 570.5 85.4 127 9 AEB7751 43 570.5 85.4 127 9 AEB7751 44 570.5 85.4 127 9 AEB7751 45 569.5 85.3 122 9 AEB7751 46 570.5 85.4 127 9 AEB7751 47 600016016017 47 600016017 48 600017 48 600017 49 600017 400016017 400016017 40	Adp22316 Human ant Adm51164 Anti-tumo Adv67314 Amino aci Adm5130 Human ant Adp22312 Human ant Adp22312 Human ant Adp22312 Human ant Adp3274 Human ant Adp36547 Single ch Ady3734 The heavy Aac35889 Human 11. Adp2128 Human ant Adp86514 Anti-RPO- Adv86514 Anti-PPO- Adv86514 Anti-PPO- Adv86514 Anti-PPO- Adv86514 Anti-PPO- Adv86514 Anti-PPO- Adv86514 Anti-PPO- Adv868175 Human ant	SNTS	Human recombinant mAb TNV148(B) heavy chain variable region. Tumour necrosis factor alpha; TNF; antibody; heavy chain; CDR; complementarity determining region; antirheumatic; antiarthritic; antiantier; antiarteriosclerotic; antiatherosclerotic; antiatherosclerotic; antiatherosclerotic; vasotropic; antianginal; cardiant; antibacterial; virucide; fungicide; antileprotic; protozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosis; therapy; TNV148; monoclonal antibody; mAb. Homo sapiens.	ide 1-7 are PCR primer-encoded and may eve sequence"	o substituted by Ser"
25 577.5 86.5 125 8 26 577 86.4 115 5 28 6.2 118 5 29 575.5 86.2 118 5 31 575.5 86.2 118 5 31 575.5 86.2 125 8 32 572.5 86.2 125 8 33 572.5 85.7 252 7 34 572.5 85.7 252 7 35 572.5 85.7 252 7 36 572 85.6 122 8 37 572.5 85.7 451 8 39 571.5 85.6 122 8 39 571.5 85.6 123 8 44 570.5 85.4 127 9 42 570.5 85.4 127 9 44 570.5 85.4 127 9 44 570.5 85.4 127 9 44 570.5 85.4 127 9 44 570.5 85.4 127 9 44 570.5 85.4 127 9 45 569.5 85.3 122 9 46 570.5 85.4 127 9 47 570.5 85.4 127 9 48 570.5 85.4 127 9 48 570.5 85.4 127 9 49 570.5 85.4 127 9 41 570.5 85.4 127 9 42 569.5 85.3 122 9 43 570.5 85.4 127 9 44 570.5 85.4 127 9 45 569.5 85.3 122 9 47 570.5 85.4 127 9 48 570.5 85.4 127 9 49 570.5 85.4 127 9 41 570.5 85.4 127 9 42 569.5 85.3 122 9 43 570.5 85.4 127 9 43 570.5 85.4 127 9 44 570.5 85.4 127 9 41 570.5 85.4 127 9 42 570.5 85.4 127 9 43 570.5 85.4 127 9 44 570.5 85.4 127 9 41 570.5 85.4 127 9 42 570.5 85.4 127 9 43 570.5 85.4 127 9 44 570.5 85.4 127 9 41 570.5 85.4 127 9 42 570.5 85.4 127 9 43 570.5 85.4 127 9 44 570.5 85.4 127 9 45 569.5 85.4 127 9 47 570.5 85.4 127 9 48 570.5 85.4 127 9 57 57 57 57 57 57 57 57 57 57 57 57 57 5	ADP2231(AAW5116, AAW5116, AAW51316, AAW51316, AD52231, AD52231, AD52231, AD52212, A		(B) heav ; TNF; 6 region; tialler erotic; acterial uroprote NV148; n	lifiers al_peptide o acids l- the native	type Pr
25 577.5 86.5 125 26 577 86.4 115 28 576 86.4 118 29 575.5 86.2 125 31 575.5 86.2 125 31 575.5 86.2 125 33 572.5 85.7 252 34 572.5 85.7 252 34 572.5 85.7 252 34 572.5 85.7 252 38 572.5 85.7 252 38 572.5 85.7 252 39 572.5 85.7 451 39 571.5 85.6 123 40 570.5 85.4 127 42 570.5 85.4 127 42 570.5 85.4 127 43 570.5 85.4 127 42 570.5 85.4 127 43 570.5 85.4 127 44 570.5 85.4 127 45 569.5 85.3 122 46 570.5 85.4 127 47 570.5 85.4 127 48 570.5 85.4 127 49 570.5 85.4 127 40 570.5 85.4 127 41 570.5 85.4 127 42 570.5 85.4 127 43 570.5 85.4 127 44 570.5 85.4 127 45 569.5 85.3 122 46 570.5 85.4 127 47 569.5 85.4 127 48 569.5 85.3 122 48 570.5 85.4 127 49 570.5 85.4 127 40 570.5 85.4 127 41 570.5 85.4 127 42 570.5 85.4 127 43 570.5 85.4 127 43 570.5 85.4 127 43 570.5 85.4 127 41 570.5 85.4 127 42 570.5 85.4 127 43 570.5 85.4 127 41 570.5 85.4 127 41 570.5 85.4 127 42 570.5 85.4 127 43 570.5 85.4 127 44 5 569.5 85.4 127 45 569.5 85.4 127 46 569.5 85.4 127 47 570.5 85.4 127 48 69100			V148 ling ing selloha ntib re	/Qua Sign amin rom	FR1 CDR1 FR2 CDR2 FR3
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The present sequence is that of the heavy chain variable region of antitumour necrosis factor (TNF) human recombinant monoclonal antibody (mAb) TWV148 [8] was produced by site-directed mutagenesis of TWV148 [8] was produced by site-directed mutagenesis of TWV148 [8] (see AAM51170), replacing residue Pro-94 with a germline Ser. TWV148 was (see AAM51170), replacing residue Pro-94 with a germline Ser. TWV148 was containing number of the man mabs produced from a GenTVV fusion using splean cells from a hybrid mouse containing human variable and constant region antibody transgenes that was immunised with human TWP alpha. The human mabs bound immobilised human TWP alpha with high avidity and had a totally human 1970; kappa isolated human, primate, rodent, ammunian, chimeric, humanised and/or complementarity determining region (CDR) -grafted anti-TWP burded and advantants, as well as anti-TWP antibody compositions, and variants, as well as anti-TWP antibody compositions, or compositions, devices, transgenic animals, transgenic composition in a cell, tissue, organ or animal (daimed) such as the matchodies may inhibit TWF induced cell adhesion molecules, inhibit TWF binding to receptor, or provide Arthritic Index improvement in a mouse model. They are useful for dispons given in the pressent celea AAM51161-6.

63). The antibodies may inhibit TWF-induced cell adhesion molecules, inhibit TWF binding to receptor, or provide Arthritic fordisp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel isolated mammalian anti-tumor necrosis factor antibody, useful for treating sickle cell anemia, diabetes, atherosclerosis, restenosis, angina pectoris, myocardial infarction, leprosy.
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29-SEP-2000; 2000US-0236826P.
01-AUG-2001; 2001US-00920137.
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99.2%;
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WO200212502-A2
                                                                          14-FEB-2002
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61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGN-YYYYGMDVWGQG 119

셤 ò 셤

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Novel isolated mammalian anti-tumor necrosis factor antibody, useful for treating sickle cell anemia, diabetes, atherosclerosis, restenosis, angina pectoris, myocardial infarction, leprosy.
                                                                                                                                                                                        complementarity determining region, antinheumatic; antiarthritic; antialcer; antiathmatic; antiallergic; antiinflammatory; antisickling; antidiabetic; antiarteriosclerotic; antiatherosclerotic; vasocropic; antianginal; cardiant; antibacterial; viruclés; fungiciés; antileprotic; protozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosis; therapy; TNV148; monoclonal antibody; mAb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence is that of the heavy chain variable region of anti-

    .19
    /label= Signal_peptide
    /note= "amino acids 1-7 are PCR primer-encoded and may differ from the native sequence"

                                                                                                                                                                                 Tumour necrosis factor alpha; TNF; antibody; heavy chain; CDR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shealy D;
                                                                                                                                                         Human recombinant mAb TNV148 heavy chain variable region.
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/label= Mature_protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'note= "encoded by A"
                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                 AAM51170 standard; protein; 146 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-AUG-2000; 2000US-0223360P.
29-SEP-2000; 2000US-0236826P.
01-AUG-2001; 2001US-00920137.
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/label= CDR3
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/label= CDR1
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/label= FR2
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/label= FR3
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/label= J6
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                                                                                                                                  10-JUN-2002 (first entry)
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N-PSDB; ABL53510.
120 TTVTVSS 126
                      140 TTVTVSS
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                                                                                                           AAM51170;
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                                                         RESULT 2
                                                                       AAM51170
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TW148. TW148 was 1 of 8 human mabs produced from a GenTNV fusion using spleen cells from a hybrid mouse containing human variable and constant region antibody transgenes that was immunised with human TWF alpha. The human mabs bound immobilised human TWF alpha with high avidity and had a totally human IgG1, kappa isotype. They showed relatedness to each other and to the human germline DP-46 heavy chain sequence (see AAM51167). The invention provides isolated human, primate, rodent, mammalian, chimeric, humanised and/or complementarity determining region (CDR)-grafted anti-TWF antibodies, immunoglobulins, cleavage products and other specified portions and variants, as well as anti-TWF antibodies, immunoglobulins, cleavage products and other specified portions of formulations, devices, transgenic animals, rand mathods of making and using them. The anti-TWF antibody compositions, formulations, devices, transgenic animals, transgenic plants, and mathods of making and using them. The anti-TWF antibody composition of making and using them. The anti-TWF antibody composition of making and using them. The anti-TWF antibody composition of making and using them of the CDRs of the composition molecules, inhibit TWF binding to receptor, or provide Arthritic Index improvement in a mouse model. They are useful for diagnosing or treating a TWF related condition in a cell anaemia, diabetes, a cardiovascular disease such a starioral arthritis, at the respection and a cardiovascular disease such as arterioral and a starioral a starioral and a such order or a cardiovascular disease such as a starioral and a starioral a starioral a starioral and a starioral and a such order or a cardiovascular disease such a starioral and a starioral a starioral a starioral a starioral a starioral a starioral and a stario
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  myocardial infarction, an infectious disease in a cell such as bacterial, viral, and fungal infections, pneumonia, leproy and malaria, a malignant disease such as leukaemia, chronic myelocytic leukaemia, Burkitt's lymphoma and multiple myeloma, or a neurological disease such as multiple sclerosis, Parkinson's disease, spinal ataxia, Alzheimer's disease and Creutzfeldt-Jakob disease. TNV148 may be especially useful as an
tumour necrosis factor (TNF) human recombinant monoclonal antibody (mAb)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNXKY
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Pred. No. 2e-48;
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Matches 125; Conservative
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119
                                                                                 80 ADSVKGRFTISRDNPKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGNYYYYYGMDVWGQG 139
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                               20 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
                                                                 61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGN-YYYYGMDVWGQG
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AAM51169 standard; protein; 146 AA AAM51169; RESULT 3 AAM51169

(first entry) 10-JUN-2002 Human recombinant mAb TNV15 heavy chain variable region.

complementarity determining region; antirheumatic; antiarthritic, antiulcer; antiasthmatic; antiallergic; antiinflammatory; antisickling; antidiabetic; antiarteriosclerotic; antiatherosclerotic; vasobropic; antianginal; cardiant; antibacterial; virucide; fungicide; antileprotic; protozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosis; therapy; TNV15; monoclonal antibody; mAb. Tumour necrosis factor alpha; TNF; antibody; heavy chain; CDR;

Homo sapiens

Location/Qualifiers Key

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The present sequence is that of the heavy chain variable region of antitumour necrosis factor (TNF) human recombinant monoclonal antibody (mAb) crumour necrosis factor (TNF) human recombinant monoclonal antibody (mAb) crumour necrosis from a hybrid mouse containing human variable and constant splen cells from a hybrid mouse containing human variable and constant region antibody transgenes that was immunised with high avidity and had a cotally human 1951, kappa isotype. They alpha with high avidity and had a cotally human 1961, kappa isotype. They showed relatedness to each other and to the human germline DP-46 heavy chain sequence (see AAMS1167). The invention provides isolated human, primate, rodent, mammalian, chimeric, humanised and/or complementarity determining region (CDR)-grafted anti-compositions and variants, as well as anti-TNF antibody. Graylementary nucleic acids, vectors, host cells, compositions, formulations, devices, transgenic animals, transgenic compositions, formulations, devices, transgenic animals, transgenic compositions, formulations and variants.

Compositions and variants a well as anti-TNF antibody, compositions, formulations of making and using them. The antibody vectors, host cells, compositions of making and using them. The antibodies may inhibit role in variable regions given in the present sequence and in AAMSI165, or ather all of the CDRs of the light chain (see AAMSI161-CC 3). The antibodies may inhibit rNF induced cell adhesion molecules, inhibit TNF binding to receptor, or provide Arthritic Index improvement in a mouse model. They are useful for diagnosing or treating a TNF relumatoid arthritis, gastric ulcer, asthma, allergic rhinitis, Crohn's pathology, sickle cell anaemia, diabetes, a caltiovascular disease such myccardial infarction, an infectious disease in a cell such as bacterial,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel isolated mammalian anti-tumor necrosis factor antibody, useful for treating sickle cell anemia, diabetes, atherosclerosis, restenosis, angina pectoris, myocardial infarction, leprosy.
1. .19
/label= Signal_peptide
/note= "amino acids 1-7 are PCR primer-encoded and may
differ from the native sequence"
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                                                                        20. .146
/label= Mature_protein
                                                                                                                                                                                                                                                                                                                                                       note= "encoded by A"
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                                                                                                                                                                                  55. .68
/label= FR2
69. .85
/label= CDR2
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01-AUG-2001; 2001US-00920137.
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                                                                                                                                                                                                                                                                                                 118. .135
/label= CDR3
                                                                                                           31. .49
/label= FR1
                                                                                                                                               50. .54
/label= CDR1
                                                                                                                                                                                                                                                                              /label= FR3
                                                                                                                                                                                                                                                                                                                                                                      136. .146
/label= J6
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07-AUG-2001; 2001WO-US024785
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viral, and fungal infections, pneumonia, leprosy and malaria, a malignant disease such as leukaemia, chronic myelocytic leukaemia, burkitt's lymphoma and multiple myeloma, or a neurological disease such as multiple eclerogis, Parkinson's disease, spinal ataxia, Alzheimer's disease and
                                                                                                                                                         61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGN-YYYYGMDVWGQG 119
                                                                                                                                                                   80 ADSVKGRFTISRDNSKNALYLQMNSLRAEDTAVYYCARDRGVSAGGNYYYYYGMDVWGQG 139
                                                                                                                   9
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                                                                                                                                                                                                                                                                                                                                           Tumour necrosis factor alpha; TNF; antibody; heavy chain; CDR; complementarity determining region; antirtheumatic; antiathritito; antialtathritito; antialtergic; antialfammatory; antisickling; antidiabetic; antiarteriosclerotic; antiatherosclerotic; vasotropic; antianginal; cardiant; antibacterial; virucide; fungicide; antileprotic; protozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosie; therapy; TNV196; monoclonal antibody; mAb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "amino acids 1-7 are PCR primer-encoded and may differ from the native sequence"
                                                                                                                   1 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
                                                                                                                              20 QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKGLBWVAFILYDGSNKKY
                                                                                                 Gaps
                                                                                                 1;
                                                                              Length 146;
                                                                                                                                                                                                                                                                                                                         Human recombinant mAb TNV196 heavy chain variable region.
                                                                                                Indels
                                                                              DB 5;
                                                                             93.9%; Score 627.5; DB 5
93.7%; Pred. No. 2.8e-46;
                                                                                                3; Mismatches
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'label= Signal_peptide
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/label= Mature_protein
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136. .146
/label= J6
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/label= CDR2
86. .117
/label= FR3
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/label= CDR1
55. .68
/label= FR2
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/label= CDR3
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/label= FR1
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                                      Creutzfeldt-Jakob disease
                                                                                                Matches 119; Conservative
                                                                                                                                                                                               TTVTVSS 126
                                                                                                                                                                                                                   TTVTVSS 146
                                                                                       Best Local Similarity
                                                          Sequence 146 AA;
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The present sequence is that of the heavy chain variable region of antitumour necrosis factor (TNF) human recombinant monoclonal antibody (mAb) TNN196 was 10 of 8 human mAbs produced from a GenTNV fusion using spleen cells from a hybrid mouse containing human variable and constant region antibody transgenes that was immunised with human TNF alpha. The human mAbs bound immobilised human TNF alpha with high avidity and had a totally human igG1, kappa isOtype. They showed relatedness to each other and to the human germline DP-46 heavy chain sequence (see AAMS1167). The invention provides isolated human, primate, rodent, mammalian, chimeric, humanised and/or complementarity determining region (CDR) grafted anti-TNF antibodies, immunoglobulins, cleavage products and other specified portions and variants, as well as anti-TNF antibody compositions, complementary nucleic acids, vectors, host cells, compositions, formulations, devices, transgenic animals, transgenic plants, and methods of making and using them. The anti-TNF antibody compositions, complementary nucleic acids, vectors, host cells, compositions, formulations of an immunoglobulin molecule, sequence and in AAMS1165, or either all of the CDRs of the heavy chain and/or light chain variable regions given in the present sequence and in AAMS1165, or either all of the CDRs of the heavy chain and/or light chain variable regions given in the present sequence and in AAMS1165, or either all of the CDRs of the heavy chain and condition in a cell, tissue, organ or animal (claimed) such as theumatoid arthritis, gastric ulcer, asthma, allergic rhinitis, crohn's characterics are accepted and parthritis, and parthritis are accepted arthritis and parthritis are accepted arthritis and parthritis and parthritis are accepted and parthritis and parthritis and parthritis and parthritis are accepted and parthritis and parthr
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4; Mismatches 4;
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                                                                                                                                                                                                                                                                     Giles-Komar J, Knight DM, Heavner G,
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07-AUG-2000; 2000US-0223360P.
29-SEP-2000; 2000US-0236826P.
01-AUG-2001; 2001US-00920137.
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92.9%;
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Creutzfeldt-Jakob disease
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Matches 118; Conservative
                                                                                                                                                                              (CENZ ) CENTOCOR INC
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RESULT 5

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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is that of the heavy chain variable region of antitumour necrosis factor (TNF) human recombinant monoclonal antibody (mAb) TYN14. TWN14 was 1 of 8 human mAbs produced from a GenTNV fusion using spleen cells from a hybrid mouse containing human variable and constant region antibody transgenes that was immunised with human TNF alpha. The human mAbs bound immobilised human TNF alpha with high avidity and had a totally human IgG1, kappa isotype. They showed relatedness to each other and to the human germline DP-46 heavy chain sequence (see AAM51167). The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated mammalian anti-tumor necrosis factor antibody, useful for treating sickle cell anemia, diabetes, atherosclerosis, restenosis, angina pectoris, myocardial infarction, leprosy.
                                                                                        complementarity determining region; antirheumatic; antiarthritic; antiulcer; antiasthmatic; antiallergic; antiinflammatory; antisickling; antidiabetic; antiarteriosclerotic; antiatherosclerotic; vasotropic; antianginal; cardiant; antibacterial; virucide; fungicide; antileprotic; protozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosis; therapy; TNV14; monoclonal antibody; mAb.
                                                                                                                                                                                                                   are PCR primer-encoded and may
                                                                                  Tumour necrosis factor alpha; TNF; antibody; heavy chain; CDR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shealy D;
                                                             Human recombinant mAb TNV14 heavy chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Scallon B,
                                                                                                                                                                                                                    /note= "amino ācids 1-7 are PCR ]
differ from the native sequence"
20. .146
/label= Mature_protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Giles-Komar J, Knight DM, Heavner G,
                                                                                                                                                                                               // label= Signal peptide
                                                                                                                                                                                                                                                                                                                                                                                                'note= "encoded by A"
                                                                                                                                                                                     Location/Qualifiers
 AAM51168 standard; protein; 146 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Fig 4; 131pp; English.
                                                                                                                                                                                                                                                                                                    55. .68
/label= FR2
69. .85
/label= CDR2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-SEP-2000; 2000US-0236826P. 01-AUG-2001; 2001US-00920137.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-AUG-2000; 2000US-0223360P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-AUG-2001; 2001WO-US024785.
                                                                                                                                                                                                                                                                                                                                            86. .117
/label= FR3
                                                                                                                                                                                                                                                                                                                                                                   118. .135
/label= CDR3
                                                                                                                                                                                                                                                                                  50. .54
/label= CDR1
                                                                                                                                                                                                                                                            31. .49
/label= FR1
                                                                                                                                                                                                                                                                                                                                                                                                          136. .146
/label= J6
                                         (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CENZ ) CENTOCOR INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-217194/27.
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                                                                                                                                                                                                                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200212502-A2
                                                                                                                                                                    Homo sapiens.
                                          10-JUN-2002
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                      AAM51168
                                                                                                                                                                                                Peptide
                                                                                                                                                                                                                                          Protein
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the mainstead and/or complementarity determining region (CDR) grafted anti-
compositions and variants, as well as anti-TWF antibody compositions,
concoding or complementary nucleic acids, vectors, host cells,
concoding or complementary nucleic acids, vectors, host cells,
compositions, formulations, devices, transgenic animals, transgenic
compositions, formulations, devices, transgenic animals, transgenic
compositions, formulations devices, transgenic animals, transgenic
compositions, formulations devices, transgenic animals, transgenic
compositions, formulations devices, transgenic animals,
compositions, formulations of makings and using them. The anti-TNF antibody
composition in and/or light chain variable regions given in the present
composition in AAMSILISS, or either all of the CDRs of the heavy chain
(see AAMSILISS-60) or all of the CDRs of the light chain (see AAMSILIS).

Composition in AAMSILISS, or either all of the CDRs of the heavy chain
composition in a cell, tissue, organ or animal (claimed) such as
condition in a cell, tissue, organ or animal (claimed) such as
arteriosclerosis, atherosclerosis, restenosis, angina pectoris or
compocardial infarction, an infectious disease in a cell such as bacterial,
compocardial infarction, an infectious disease in a cell such as bacterial,
compocardial infarctions, pneumonia, leprosy and malaria, a malignant
compocardial infarctions, or a neurological disease such as multiple
compocardial disperses or a neurological disease and a malignant or animal compocardial disperse and and malaria, a malignant or all and published to the compocardial disperse and an early and the compocardial disperse and an entry or animal compoca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGN-YYYYGMDVWGQG 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  79
provides isolated human, primate, rodent, mammalian, chimeric,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sclerosis, Parkinson's disease, spinal ataxia, Alzheimer's disease and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human heterodimeric antibody; human; antibody; binding affinity; protective antigen; Bacillus anthracis; anthrax infection; cell recept edema factor; lethal factor; virucide; antibacterial; immunotherapy; anti-toxin; anti-infective; anthrax; botulinum; smallpox; Venezuelan equine encephalomyelitis virus; VEEV; West Nile virus; WNV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGKGLEWVAILLYDGSSKKY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human heterodimeric antibody heavy chain variable region SEQ ID NO:98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 615.5; DB
Pred. No. 3e-45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2002US-0376408P.
2002US-0414053P.
2002US-0428807P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92.1%;
92.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Creutzfeldt-Jakob disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 92.9
Matches 118; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TTVTVSS 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  140 Triviryss 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 146 AA;
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27-SEP-2002;
25-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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27-SEP-2002; 2002US-0414053P.
25-NOV-2002; 2002US-0428807P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GTTVTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 135 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO2004110362-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-DEC-2004.
                                                                                                     Bowdish KS,
                                                                                                                                                                                                                                                   Nile virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADV86823;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         123
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  g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ઠે
                                                                                                                                                                                                                                                                                    The present invention describes a human heterodimeric antibody (I) (fragment) having a binding affinity of at least 1x10-8 M to the protective antigen of Bacillus anthracis or a molecule involved in anthrax infection that blocks binding of the antigen or molecule to receptors, edema factor and lethal factor. (I) has virucide and antibacterial activities, and can be used in immunotherapy. The antibodies (I) are useful as anti-toxins or anti-infectives with respect to infective agents, such as anthrax, botulinum, smallpox, Venezuelan equine encephalomyelitis virus (VEBV), or West Nile virus (WND). The present sequence represents a human heterodimeric antibody heavy chain variable region amino acid sequence, which is used in the exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ï
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                122
                                                                                                                                          New human heterodimeric antibodies or their antibody fragments, useful as anti-toxins or anti-infectives with respect to infective agents, e.g. anthrax, botulinum, smallpox, Venezuelan equine encephalomyelitis or West
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADSVKGRFTISRDNSKNTLYLQMINSLRAEDTAVYYCARDRGIAAGGN--YYYYGMDVWGQ 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human heterodimeric antibody; human; antibody; binding affinity; protective antigen; Bacillus anthracis; anthrax infection; cell recept edems factor; lethal factor; virucide; antibacterial; immunotherapy; anti-toxin; anti-infective; anthrax; botulinum; smallpox; venezuelan equine encephalomyelitis virus; VEEV; West Nile virus; WNV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human heterodimeric antibody heavy chain variable region SEQ ID NO:14.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 135;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        87.3%; Score 583; DB 7;
86.7%; Pred. No. 1.7e-42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4; Mismatches
                                                                                                                                                                                                                                                 Claim 11; SEQ ID NO 98; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADD28236 standard; protein; 135 AA
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2002US-0376408P.
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                     (ALEX-) ALEXION PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 111; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GTTVTVS 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  130
                                                                                                   WPI; 2003-722327/68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 135 AA;
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29-APR-2002;
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                                                              KS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JAN-2004
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                                                                                                                                                                                                          Nile virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  123
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                                                            Bowdish
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ID ADD2

XX ADD2

XXX ADD2

XXX IS-0.

XXX IS-0.

XXX Hums

XXX Hums

XXX ADD2

XXX AD
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The present invention describes a human heterodimeric antibody (I) (fragment) having a binding affinity of at least IX10-8 M to the protective antigen of Bacillus anthracis or a molecule involved in anthrax infection that blocks binding of the antigen or molecule to receptors, edema factor and lethal factor. (I) has virucide and antibacterial activities, and can be used in immunotherapy. The antibodies (I) are useful as anti-toxins or anti-infectives with respect to infective agents, such as anthrax, botulinum, smallpox, Venezuelan equine encephalomyelitis virus (VEEV), or West Nile virus (WND). The present sequence represents a human heterodimeric antibody heavy chain variable region amino acid sequence, which is used in the exemplification
                                                                                                                                                                                             New human heterodimeric antibodies or their antibody fragments, useful as anti-toxins or anti-infectives with respect to infective agents, e.g. anthrax, botulinum, smallpox, Venezuelan equine encephalomyelitis or West
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 135;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 583; DB 7;
Pred. No. 1.7e-42;
4; Mismatches 11.
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                                                                                                                                                                                                                                                                                                                                                                        Claim 6; SEQ ID NO 14; 67pp; English.
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86.7%;
(ALEX-) ALEXION PHARM INC.
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                                                                                                                                     WPI; 2003-722327/68.
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comprises administering an antibody or antibody fragment that binds to a comprises administering an antibody or antibody fragment that binds to a molecule involved in anthrax infection and which has the ability to block the binding of the molecule to at least one of a cell receptor, PA63, edecamining exposure to B. antiracis comprises assaying a sample for the presence of a molecule selected from cell receptors, PA63, PA63 heptamer, PA83, edema factor or lethal factor with an antibody that has binding a faffinity for the molecule, where the presence of elevated levels of the antibody correlates with the presence of elevated levels of the cantibody to a cell receptor. PA63, PA63, PA63 heptamer, PA83, edema factor or lethal factor with a secondary antibody having binding affinity for the antibody, where the presence of elevated levels of the scondary antibody correlates with the presence of elevated levels of the scondary antibody correlates with the presence of elevated levels of the secondary antibody correlates with the presence of Elevated levels of the secondary antibody correlates with the presence of Elevated levels of the secondary antibody correlates with the presence of Elevated levels of the secondary antibody and a light chain sequence of Elevated levels of the secondary antibody and a light chain lambda region selected from a group of sequences ADV86815 and a light chain lambda region selected from a group of sequences ADV86815. The second sequences ADV86815 and a light chain lambda region selected from a group of sequences ADV86815. The sesent invention are also useful for producing anticoming or anti-infectives to infective agents such as antirax, botulinum, smallpox, venezuelan equine encephalomyelitis and west Nile virus. The compositions of the present invention are also useful for producing and the sequence is that of the heavy chain variable region elected from a group of sequences and pA83. Phage libraries were developed from mand padd and pA83. Phage libraries padd and pA83. Phage libraries 
                                                                                                                                                                                 Treating an animal with anthrax infection by administering an antibody to the protective antigen of Bacillus anthracis, and blocking binding to cell receptors, edema factor or lethal factor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    against anthrax. The libraries were panned against PA83 and PA63, and sequence analysis was performed on positive responders. Neutralization of anthrax toxin activity by purified Fabs was demonstrated.
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                                                                                         Nolan MJ
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                                                                                         Maruyama T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               87.3%; Score 583; DB 9;
86.7%; Pred. No. 1.7e-42;
iive 4; Mismatches 11;
                                                                                         Wild MA,
                                                                                                                                                                                                                                                                             Claim 7; SEQ ID NO 14; 87pp; English.
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                                                                                         Frederickson S,
02-JUN-2003; 2003US-00452593
                                           (ALEX-) ALEXION PHARM INC.
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                                                                                         Bowdish KS,
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The present invention describes a protein comprising an immunoglobulin (1g) heavy chain variable (VH) domain and an Ig light chain variable (VL) domain. The protein binds a complex comprising a major histocompatibility complex (MHC) and a peptide, does not substantially bind the MHC in the absence of the bound peptide, and does not substantially bind the peptide in the absence of the MHC. The peptide is a peptide fragment of gp100, MUC1, TAX or hTERT. Also described: (1) a pharmaceutical composition comprising the novel protein and a carrier; (2) a cyctoxic T cell comprising one or more nucleic acids for expressing the Ig that binds a comprising an MHC and a peptide, does not substantially bind the peptide in the absence of the bound peptide, and does not substantially bind the comprising a first segment that encodes the Ig variable domain; (4) a host cell comprising hererologous nucleic acid sequences that encodes the protein; (5) a transgenic animal whose genome includes heterologous nucleic acid sequences that encode the protein; (6) identifying the protein that specifically binds the MHC. Peptide complex; (7) expressing an antigen-binding protein; (8) abbating or killing a target cell that displays a peptide on a surface MHC molecule; (9) treating or preventing a concerous disorder in a subject; and (10) detecting an MHC-peptide composition for treating or preventing a composition be needed to the new of the invention has cytostatic and MHC-peptide composition for treating or preventing a composition for treating or preventing a composition for preventing a composition for the period of an antibod which because the provention and composition for the period of an antibod which because the period of an antibod of an antibod of an antibod of a period of a period of an antibod of an antibod of an antibod of a 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New protein comprising an immunoglobulin heavy chain variable (VH) domain and an immunoglobulin light chain variable (VL) domain, useful for preparing a composition for treating or preventing a cancerous disorder.
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                                              immunoglobulin; Ig; heavy chain variable domain;
light chain variable domain; major histocompatibility complex; MHC;
gp100; MUC1; TAX; hTERT; cytostatic; gene therapy; cancerous disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 QVQLVQSGGGVVQPGRSLRLSCAASGFTFSSYGMHWYRQAPGKGLEWVAVISYDGSNKYY
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Human antibody T3F2 heavy chain amino acid sequence SEQ ID NO:82.
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88.9%; Pred. No. 2e-42;
ive 3; Mismatches 9; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (TECR ) TECHNION RES & DEV FOUND LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                       20-FEB-2002; 2002US-0358994P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 112; Conservative
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                                                                                                                                                                                                      Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a method of identifying binding site domains (BSD) that retain the capacity of binding to a predetermined epitope when positioned C-terminal of at least one further domain in a recombinant bicor multivalent polypeptide. The method comprises (a) testing a panel of the SDD displayed on the surface of a blological display system as part of the subject of the surface of a blological display system as part of the fusion protein for binding to a predetermined epitope, where the fusion protein comprises an additional domain positioned N-terminal of the BSD and an amino acid sequence that mediates anchoring of the fusion protein to the surface of the display system; and (b) identifying a BSD that to binds to the predetermined epitope. The method is useful to identify bicor multivalent polypeptides that comprise antibody binding sites capable of efficiently binding to the corresponding antigen. The polypeptides or antibodies identified by the method are useful therapeutically and antibodies identified by the method are useful therapeutically and flagments that bind independently of their position within bifunctional single-chain fusion proteins can be isolated from combinatorial antibody libraries using the new in vitro method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Phage display system for identification of binding site domains retaining capacity to bind an epitope.
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                                                                                                                                                                                                           Binding site domain; BSD; epitope; fusion protein; therapeutic; cancer; autoimmune disease; scFv-antibody; single-chain Fv.
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Pred. No. 2.2e-42;
5; Mismatches 10
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                                                                                                                                                                                 Human D4.5 heavy chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Fig 3.1; 152pp; English.
                                                                                             AAY17954 standard; protein; 127 AA
                                                                                                                                                                                                                                                                                                                                                                                                                      Borschert K,
                                                                                                                                                                                                                                                                                                                                     98WO-EP007313.
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87.4%;
                                                                                                                                                     (first entry)
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Best Local Similarity 87.4
Matches 111; Conservative
121 TVTVSS 126
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                         119 TVTVSS 124
                                                                                                                                                                                                                                                                                                                                                                                                                      Raum T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB; AAX77236.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 127 AA;
                                                                                                                                                                                                                                                                                                                                                                                          KUFE/) KUFER
                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                             WO9925818-A1
                                                                                                                                                                                                                                                                                                                                                               17-NOV-1997;
                                                                                                                                                    04-AUG-1999
                                                                                                                                                                                                                                                                                                          27-MAY-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                    Kufer P,
                                                                                                                          AAY17954;
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This is the amino acid sequence of the human D4.5 heavy chain variable region, used in the method of the invention. for providing receptors that can be used for targeting antigens in humans without being immunogenic themselves. Such receptors can be used for treating diseases such as tumours or auto-immune diseases, graft rejection after transplantation, infectious diseases by targeting cellular receptors as well as allergic, inflammatory, endocrine and degenerative diseases by targeting key molecules involved in the pathological process
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Production of anti-human antigen receptors - by selecting a combination of functionally rearranged VH and VL immunoglobulin chains expressed from a recombinant vector.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61 ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARDRGIAAGGN-YYYYGMDVWGQG 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 EVQLLESGGGVVQPGRSLRLSCAASGFTFSSYGMIMVRQAPGKGLEWVAVISYDGSNKYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
                                                                                                                                                                                                                                                              Human; D4.5 heavy chain variable region; receptor; antigen; tumour; auto-immune disease; graft rejection; allergy; inflammatory disease; endocrine disease; degenerative disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1;
                                                                                                                                                                                                              Amino acid sequence of human D4.5 heavy chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 138;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 581.5; DB 2;
Pred. No. 2.4e-42;
5; Mismatches 10;
                                               AAW80815 standard; protein; 138 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AEC16143 standard; protein; 457 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 9; Fig 7; 84pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98WO-EP002180.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97EP-00106109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      87.48;
                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kufer P, Raum T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-PSDB; AAV68537.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (KUFE/) KUFER P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 138 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RAUM T.
                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9846645-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-0CT-1998
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                                                                                                       AAW80815;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RAUM/)
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AEC16143
ID AEC16
RESULT 11
                           AAW80815
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120 TTVTVSS 126

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The invention relates to a method of treating tumorous disease by administering a human immunoglobulin specifically binding to the human epithelial cell adhesion molecule (EpCAM) also termed as 17-1A antigen, KSA, EGP40, GA733-2, ks 1-4 and esal antigen. The immunoglobulin exhibits a serum half-life of at least 15 days and is administered no more composition comprising anti-EpCAM are useful for treating tumorous disease, such as breast cancer, epithelial cancer, hepatocellular carcinoma, cholangiocellular cancer, stomach cancer (melanoma), a cancer of the urogenital tract (e.g., ovarian cancer, endometrial cancer, carcin cancer, and kidney cancer), lung cancer, gastric cancer, a cancer of the small intestine, liver cancer, pancreas cancer, and cancer of the cancer of the bile duct, esophagus cancer, gall bladder cancer, a cancer of the thyroid gland. The present sequence is the human anti-EpCAM immunoglobulin heavy chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Treating tumorous disease, such as breast, colon, prostate, liver, skin, ovarian, cervical and lung cancer, by administering a human immunoglobulin specifically binding to the human epithelial cell adhesion molecule (EpcAM) antigen.
                                                                                                                                       pancreas tumor;
                                                                                                                                 Antibody therapy; tumor; cancer; cytostatic; neoplasm; pancreas tumor; ovary tumor; thyroid tumor; stomach tumor; breast tumor; epithelial cancer; hepatocellular carcinoma; cholangiocellular tumor; head & neck tumor; win tumor; melanoma; uterine cancer; uterine cervix tumor; renal tumor; lung tumor; intestine tumor; liver tumor; gallbladder carcinoma; esophagus tumor; immunoglobulin; antibody; epithelial cell adhesion molecule; heavy chain.
                                                                                                Human anti-EpCAM immunoglobulin heavy chain, SEQ ID NO: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Quadt C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 12; SEQ ID NO 1; 23pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prang N,
                                                                                                                                                                                                                                                                                                                                                                                                                  13-FEB-2004; 2004US-00778915.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-FEB-2004; 2004US-00778915.
                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peters M, Locher M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2005-590351/60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MICR-) MICROMET AG.
                                                                                                                                                                                                                                                                                                                                      US2005180979-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 457 AA;
                                                                                                                                                                                                                                                                                                 Homo варіепв
                                                         20-OCT-2005
                                                                                                                                                                                                                                                                                                                                                                             18-AUG-2005
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ä 1 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY Gaps ä DB 9; Length 457; 10; Indels 87.1%; Score 581.5; DB 9 87.4%; Pred. No. 8.1e-42; ive 5; Mismatches 10 Best Local Similarity 87.4 Matches 111; Conservative Query Match 셤 Š

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epithelial cell adhesion molecule; inflammatory cytokine; GM-CSF; granulocyte/macrophage colony stimulating factor; heterominibody; granulocyte/macrophage colony stimulating factor; heterominibody; immunostine; multipunding constant domain; immunostiopulin; cytostantc; immunostimulatory; antileukaemia; diagnosis; antiproliferative; prevention; treatment; malignant; haematopoletic cell;
                                                                                             Epcam;
                                                                                                                                                                                                                                                                                                                                                                        New multifunctional compounds useful for preventing and/or treating malignant cell growth and for detection and diagnosis.
                                                                                             scPv; 17-1A antigen; human;
                                                                                                                                                              lymphoma; leukaemia; solid tumour; carcinoma; melanoma; sarcoma
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                                                                                                                                                                                                                                                                                                                     Zett1
                                                                                                                                                                                                                                                                                                                     Borschert K,
                                                                                                                                                                                                                                                                                               (MICR-) MICROMET GES BIOMEDIZINISCHE FORSCHUNG.
                                                                                             single-chain variable fragment;
                                                                                                                                                                                                                                                                                                                                                                                                            Example 10; Fig 55A; 166pp; English.
         AAY44994 standard; protein; 523 AA
                                                                                                                                                                                                                                                                                                                     Dreier T, Baeuerle PA,
                                                                                                                                                                                                                                                    99WO-EP005416.
                                                                                                                                                                                                                                                                           98EP-00114082.
                                                                        HD70scFv-CH1-GM-CSF chain.
                                                                                                                                                                                                                                                                                                                                           WPI; 2000-195265/17.
                                                                                                                                                                                                                                                                                                                                                        N-PSDB; AAZ50587
                                                                                                                                                                                                          WO200006605-A2.
                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                     28-JUL-1999;
                                                                                                                                                                                                                                                                           28-JUL-1998;
                                                  23-MAY-2000
                                                                                                                                                                                                                                10-FEB-2000.
                               AAY44994;
                                                                                                                                                                                                                                                                                                                     Kufer P,
AAY44994
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The patent discloses heterominibodies which are multifunctional compounds produceable in a mammalian host cell as a secretable and fully functional thererodiner of two polypeptide chains, where one of the polypeptide chains chains comprises, a CHI-domain (constant domain of an immunoglobulin chain) and the other chain (constant domain of an immunoglobulin light chain). The polypeptide chains further comprise, can immunoglobulin light chain). The polypeptide shaving chain the constant domains at least two (poly) peptides having different receptor or ligand functions, where further at least two of the different (poly) peptides lack an intrinsic affinity for one another and are linked via the constant domains. The heterominbodies have cytostatic, immunostimulatory, antileukaemia and antiproliferative cytostatic, immunostimulatory, antileukaemia and antiproliferative cytostating malignant cell growth related to malignancies of haematopoietic cells e.g. lymphomas and leukaemias, or to solid tumours e.g. carcinomas, cellance of present sequence is the left chain of an elanomas and sarcomas. The present sequence is the left chain of an elanoma and sarcomas. The present sequence is the left chain of the certain inflammatory cytokine granulocyte/macrophage colony stimulating factor (GM-CSF), plus a hexahistidine sequence for ease of purification. CHD70 screption (CHD70 screption) antigen

Sequence 523 AA;

9 1 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY Gaps 1; Length 523; Indels 87.1%; Score 581.5; DB 3; 87.4%; Pred. No. 9.3e-42; ive 5; Mismatches 10; Matches 111; Conservative Similarity Query Match Local

ò 셤 9.3e-42;

Pred. No.

87.48;

Best Local Similarity

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produceable in a mammalian host cell as a secretable and fully functional beterodimer of two polypeptide chains, where one of the polypeptide chains comprises a CLT-domain (constraint domain of an immunoglobulin heavy chain) and the other chain comprises CL-domain of an immunoglobulin hight chain). The polypeptide chains further comprise, can immunoglobulin light chain). The polypeptide chains further comprise, clused to the constant domains at least two (poly)peptides having different receptor or ligand functions, where further at least two of the different receptor or ligand functions, where further at least two of the clifferent receptor or ligand functions, where further at least two of the different receptor or ligand functions, where further at least two of the clifferent (poly)peptides lack an intrinsic affinity for one another and are linked via the constant domains. The heterominibodies have cytostatic, immunostimulatory, antileukaemia and antiproliferative activities. These compounds can be used for diagnosing, preventing and treating malignant cell growth related to malignancies of haematopoietic cells e.g. lymphomas and leukaemias, or to solid tumours e.g. carcinomas, melanomas and sarcomas. The present sequence is the right chain of a heterominibody comprising HD70 single-chain Fv (scFv) fragment N-cerminally linked to human Ck domain (constant domain of immunoglobulin-kappa light chain) which bears at its Cterminus the human inflammatory cytokine interleukin-2 (III-2). HD70 scFv specifically recognises the human epithelial cell adhesion molecule (BpCAM) also called 17-1A antigen
ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGN-YYYYGMDVWGQG 119
                      ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKDMGWGSGWRPYYYYGMDVWGQG 261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 patent discloses heterominibodies which are multifunctional compounds
                                                                                                                                                                                                                                                                                                                                                                                                             HD70; single-chain FV fragment; scFV; antibody; 17-1A antigen; human; bpCAW; epithellal cell adhesion molecule; inflammatory cytckine; IL-2; interleukin-2; CK-domain; kappa light chain constant domain; heterominibody; multifunctional compound; immunoglobulin; cytostatic; immunostimulatory; antileukaemia; diagnosis; prevention; antiproliferative; treatment; malignant; haematopoietic cell; lymphoma; leukaemia; solid tumour; carcinoma; melanoma; arcoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         compounds useful for preventing and/or treating the and for detection and diagnosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MICR-) MICROMET GES BIOMEDIZINISCHE FORSCHUNG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 10; Fig 55B; 166pp; English.
                                                                                                                                                                                                                                       AAY44995 standard; protein; 524 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Baeuerle PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  malignant cell growth and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99WO-EP005416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98EP-00114082
                                                                                                                                                                                                                                                                                                                                                                   HD70scFv-Ck-interleukin 2.
                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                   TTVTVSS 126
                                                                                                                             268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           multifunctional
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                                                                                                                             TTVTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-195265/
N-PSDB; AAZ50588.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200006605-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 524
                                                                                                                                                                                                                                                                                                                          23-MAY-2000
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61
                                        202
                                                                                   120
                                                                                                                                                                                                                                                                                  AAY44995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New
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DB 3; Length 524;

87.1%; Score 581.5;

Query Match

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                                                            201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a human monoclonal antibody that binds to platelet-derived growth factor-D (PDGF-D). The antibodies are useful for recating chronic and recurrent human diseases, such as inflammation, autoimmunity and cancer. The PDGF-D nucleic acids and polypeptides are useful for modulating collagen formation, and for staging various cancers. Antibodies to platelet-derived growth factor-D (PDGF-D) were generated using an active protein fragment of the gene product from the clone 30664188.0.99 arising in the conditioned medium obtained when HEKZ93 cells are transfected with the plasmid pCEP4/Sec-30664188. This sequence corresponds to a protein used in the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9
                               9
                                                                                                           202 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKDMGWGSGWRPYYYYGMDVWGQG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  142 EVQLLESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYDGSNKYY
                                                                                           61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGN-YYYYGMDVWGQG
                             QVQLVESGGGVVQPGRSLRLSCAASGPIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              growth
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             factor-D (PDGF-D), useful for treating chronic and recurrent diseases, such as inflammation, autoimmunity and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                platelet-derived
                                                                                                                                                                                                                                                                                                                                                                                              antiinflammatory; immunomodulator; cytostatic; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  87.0%; Score 581; DB 7; Length 126; 88.1%; Pred. No. 2.4e-42; ive 3; Mismatches 12; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gazit G,
 Indels
                                                                                                                                                                                                                                                                                                                                                               Anti-human PDGF-D antibody heavy chain protein sequence.
10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chen F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New human monoclonal antibody that binds to
 5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 17; 255pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yang X,
                                                                                                                                                                                                                                                                    ADK18593 standard; protein; 126 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Feng X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-JAN-2003; 2003WO-US000398.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JAN-2002; 2002US-00041860
                                                                                                                                                                                                                                                                                                                                  06-MAY-2004 (first entry)
 Matches 111, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 88.1
Matches 111; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jia X,
                                                                                                                                                                                       268
                                                                                                                                                         TTVTVSS 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-587119/55.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ABGE-) ABGENIX INC.
                                                                                                                                                                                        TTVTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 126 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003057857-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Corvalan JRF,
                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
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121 TVTVSS 126 |||||| 121 TVTVSS 126

Search completed: December 28, 2005, 13:58:06 Job time : 149.538 secs

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries Database

PIR 80:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Ig heavy chain V r Description SUMMARIES \$48797 \$338493 \$31106 \$31106 \$31106 \$449028 \$449028 \$31679 \$31177 \$31177 \$31177 \$31177 \$31177 \$31177 \$31177 \$31609 \$31609 \$31609 \$31609 \$31609 \$31609 PH1645 S46391 S36270 E36005 B Length & Query Match 1 577 563.5 539.5 539.5 534.5 534.5 535.5 536.5 528.5 52 Result

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133	109	109	123	118	126	145	113	151	122	108	119	121	125	111	127
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ALIGNMENTS

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Total number of hits satisfying chosen parameters:

seq length: 0 seq length: 200000000

Minimum DB Maximum DB

RESULT 1 E36005 Igheavy chain V region (M72) - human C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Species: Homo sapiens (man) C;Date: 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 16-Dec-1998 C;Accession: E36005 R;Schroeder Jr., H.W.; Wang, J.Y. Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990 R;Schroeder Jr., H.W.; Wang, J.Y. Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990 A;Atitle: Preferential utilization of conserved immunoglobulin heavy chain variable gene A;Accession: E36005 A;Accession: E36005 A;Accession: E36005 A;Accession: B26005 A;Accession: B26005 A;Cross-references: UNIPARC:UPI0000176C30; GB:M34030 C;Genetics C;Genetics A;Cross-references: GDB:118731; CMIM:146910 A;Cross-references: GDB:118731; CMIM:146910 A;Cross-references: GDB:118731; Mimunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology < IMM>	Query Match 86.4%; Score 577; DB 2; Length 122; Best Local Similarity 88.3%; Pred. No. 6.9e-45; Matches 112; Conservative 4; Mismatches 6; Indels 4; Gaps 1; QY 1 QVQLVBSGGGVVQPGRSLELSCAASGFIFSSYAMHWVRQAPGMGLEWVAPRSYDGSNKKY 60 QY 61 ADSVKGRFITSRDNSKNTLXLQMNSLRAEDTAVYYCARDRGLBWAVISYDGSNKYY 60 QY 61 ADSVKGRFITSRDNSKNTLXLQMNSLRAEDTAVYYCARDRGLAAGGNYYYYGMDVWGQGT 120 QY 61 ADSVKGRFITSRDNSKNTLXLQMNSLRAEDTAVYYCARDRHSSSWYYGMDVWGQGT 116 Db 61 ADSVKGRFITSRDNSKNTLXLQMNSLRAEDTAVYYCARDRHSSSWYYGMDVWGQGT 116 QY 121 TVTVSS 126 Db 117 TVTVSS 122 CS 117 TVTVSS 122 ASSTANT AND	A;Molecule type: mRNA
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Ig heavy chain - human
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens
C;Date: 0.2-Dec-1993 #sequence_revision 26-May-1995 #text_change 31-Dec-2004
C;Accession: 531116
R;Raaphorst, F.M.; Timmers, B.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman R;Raaphorst, F.M.; Timmers, B.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman A;Reference number: 831104; MUID:92111633; PMID:1730252
A;Accession: S31116
A;Accession: S31116
A;Scatus: preliminary; nucleic acid sequence not shown; translation not shown
A;Residues: 1-118 cRAA.
R;Schroeder Jr., H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin heavy chain variable gene A;Reference number: A36005; MUID:90349571; PMID:2117273
A;Accession: F36005
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A; Once: the nucleotide sequence was submitted to the EMBL Data Library, October 1991
C; Superfamily: immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61 ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARDRGIAAGGNYYYYGMDVWGQGT 120
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84.1%; Pred. No. 1.5e-41;
ive 3; Mismatches 10
                                                                                                                                                                                                                                                                                                                                                         A;Gene: GDB:IGH0; IGHDY1
A;Cross-references: GDB:118731; OMIM:146910
A;Map position: 14q32.33-14q32.33
C;Superfamily: immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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Matches 106; Conservative
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A;Molecule type: mRNA
A;Residues: 1-119 <SCH>
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Best Local (
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   A;Residues: 1-128 <MAH>
A;Cross-references: UNIPARC:UPI0000116700; EMBL:Z46379; NID:g587147; PIDN:CAA86512.1; PI B;Tomlinson, I.M.; Walter, G.; Marks, J.D.; Llewelyn, M.B.; Winter, G. Di., Tomlinson, I.M.; Walter, G.; Marks, J.D.; Llewelyn, M.B.; Winter, G. Di., 776-798; MID:93021117; PMID:1404388
A;Reference number: S26885; MUID:93021117; PMID:1404388
A;Reference number: S26885; MUID:93021117; PMID:1404388
A;Residues: 1-98 <TOM>
A;Residues: 1-98 <TOM>
C;Superferences: UNIPARC:UPI0000038183; EMBL:Z12350; NID:g32922; PIDN:CAA78220.1; PID C;Superferences: Universe immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology <IMM>
F;15-98/Domain: immunoglobulin homology <IMM>
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19 heavy chain - human (fragment)
C;Species: Homo sapiens (man)
C;Accession: 638493
R;Marks, J.D.; Ouwehand, W.H.; Bye, J.M.; Finnern, R.; Gorick, B.D.; Voak, D.; Thorpe, E submitted to the EMBL Data Library, June 1993
A;Bescription: Human antibody fragments specific for human blood group antigens from a falserance number: 838488
A;Accession: 638493
A;Accession: 638493
A;Accession: 538488
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F36005
IG heavy chain V region (M49) - human
IG heavy chain V region (man)
C;Species: Homo sapiens (man)
C;Date: 21.Dec.1990 #sequence_revision 21-Dec-1990 #text_change 31-Dec-2004
C;Accession: F36005
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Pred. No. 1.1e-44;
3; Mismatches 11; Indels
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87.3%; Pred. No. 1.1e-43;
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Best Local Similarity 87.5%;
Matches 112; Conservative
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Matches 110; Conservative
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C;Accession: A49028
R;Timmers, B.; Kenter, M.; Thompson, A.; Kraakman, M.B.; Вектап, J.B.; Alt, F.W.; Schuur
B;Timmers, B.; Kenter, M.; Thompson, A.; Kraakman, M.B.; Вектап, J.B.; Alt, F.W.; Schuur
B;Title: Diversity of immunoglobulin heavy chain gene segment rearrangement in B lymphoc
A;Reference number: A49028; MUID:92008140; PMID:1915549
                                                                                                                                                                                                                                                                                                                   A; Accession: A49028
A; Status: preliminary
A; Molecule type: mRNA
A; Molecule type: manuaglobulin to MOBI backbone (NCBIN: 64471, NCBIP: 64470)
C; Superfamily: immunoglobulin to manuaglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology < NMM>
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                                                                        Ig heavy chain V-III region - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 21-Jan-1994 #sequence_revision 18-Nov-1994 #text_change 23-Jul-1999
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Pred. No. 6.9e-41;
2; Mismatches 7; Indels
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83.6%; Pred. No. 7.3e-41;
iive 2; Mismatches 12
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A;Molecule type: DNA
A;Residues: 1-114 <FIG>
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Matches 105;
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Ig heavy chain V region (M74) - human
C;Species: Homo sapiens (man)
C;Date: 21-Dec-1900 #sequence_revision 21-Dec-1990 #text_change 31-Dec-2004
C;Accession: G36005
R;Schroeder Jr., H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin heavy chain variable gene A;Reference number: A36005; MUID:90349571; PMID:2117273
A;Accession: G36005
A;Residues: Preliminary
A;Molecule type: mRNA
A;Residues: 1-121 <SCH>
A;Cross-references: UNIPROT:Q8WUXL; UNIPARC:UPI0000176C2C; GB:M34031
C;Genetics: GB:IGHDY1
A;Cross-references: GDB:118731; OMIM:146910
A;Gene GBB:IGHOXI = 14432.33
C;Superfeamily: immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
P;15-98/Domain: immunoglobulin homology <IMM>
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Pred. No. 4.4e-41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4; Mismatches
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Local Similarity 84.1%;
nes 106; Conservative 4
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Best Local Simi
Matches 106;
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Ig heavy chain V region - human (fragment)
C;Species: Homo sapiens (man)
C;Accession: 831679
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
Submitted to the EMBL Data Library, June 1992
A;Description: Mechanisms that generate human immunoglobulin diversity operate from the A;Reference number: 831585
A;Accession: 831679
A;Accession: 831679
A;Accession: L14 «CUI»
A;Residues: 1-134 «CUI»
A;Residues: 1-134 «CUI»
A;Coss-references: UNIPARC:UPI0000116475; EMBL:Z14203; NID:g30965; PIDN:CAA78572.1; PIE
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology «IMM»
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C;Species: Homo sapiens (man)
C;Date: 24-Jul-1998 #sequence_revision 24-Jul-1998 #text_change 31-Dec-2004
C;Accession: S70442
Mol. Immunol. 29, 1363-1373, 1992
A;Title: IgM kappa/lambda BBV human B cell clone: an early step of different A;Reference number: S70442; MUID: 93024508; PMID:1383695
A;Reference number: Growth Conceptual translation
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C;Superfamily: immunoglobulin homology
F;34-117/Domain: immunoglobulin homology <IMM>
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                                                                                                                                                                                                                                                                                                                                                                                                     79.1%; Score 528.5; DB 2.83.5%; Pred. No. 1.7e-40; tive 2; Mismatches 6
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Best Local Similarity 83.55
Matches 106; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: mRNA
A; Residues: 1-140 <CUI>
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Best Local Similarity
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S46392
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S1966
Ig heavy chain V region (VH3DJH4) - human
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Species: 10566
C;Accession: S19666
R;Marks, J.D.; Hoogenboom, H.R.; Bonnert, T.P.; McCafferty, J.; Griffiths, A.D.; Winter, J. Mol. Biol. 222, 581-597, 1991
A;Title: By-passing immunization. Human antibodies from V-gene libraries displayed on ph A;Recession: S19666
A;Molecule type: mRNA
A;Accession: S19666
A;Molecule type: mRNA
A;Cross-references: UNIPARC:UPIO000115FE5; EMBL:X61646; NID:g37688; PIDN:CAA43827.1; PIC
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
  RESULT 10
831603
Ig heavy chain V region - human
C; Species: Homo sapiens (man)
C; Date: 03-Mar-1994 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C; Accession: 331603
R; Cuitsinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
submitted to the EMBL Data Library, June 1992
R; Cuitsinier, A.M.; Gauthiers that generate human immunoglobulin diversity operate from the A; Reference number: 831603
A; Accession: 831603
A; Residues: preliminary
A; Molecule type: mRNA
A; Residues: 1-132 <CUI->
A; Cross-references: UNIPARC:UP10000116455; EMBL; Z14168; NID:930999; PIDN:CAA78537.1; PIC
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotesramer; immunoglobulin
F; 30-113/ Domain: immunoglobulin homology <IMM>
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82.5%; Pred. No. 1.2e-40;
iive 6; Mismatches 11.
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Pred. No. 1.1e-40;
4; Mismatches 9
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82.5%;
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Best Local Similarity 82.5
Matches 104; Conservative
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Ig heavy chain - human

Ig heavy chain - human

Ig heavy chain - human

C; Species: Home saplens (man)

C; Species: Home saplens (man)

C; Accession: S31117

R; Raaphorst, F. M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurman Eur., J. Immunol. 22, 247-251, 1992

A; Title: Restricted utilization of germ-line V(H)3 genes and short diverse third complem A; Recence number: S31104; MUID:9211633; PMID:1730252

A; Accession: S31117

A; Status: preliminary; nucleic acid sequence not shown; translation not shown

A; Status: preliminary; nucleic acid sequence not shown; translation not shown

A; Stesidues: 1-122 - RAA>

A; Cross-references: UNIPARC:UPI0000176C8D; EMBL:X62967

A; Note: the nucleotide sequence was submitted to the EMBL Data Library, October 1991

C; Superfamily: immunoglobulin V region; immunoglobulin homology

C; Keywords: heterotetramer; immunoglobulin

F; 15-98/Domain: immunoglobulin homology < INM>
C;Species: Homo sapiens (man)
C;Date: 27-Jan-1995 #text_change 20-Jun-2000
C;Accession: 846392
R;Pigini, M.; Marks, J.D.; Winter, G.; Griffiths, A.D.
J. Mol. Biol. 239, 68-78, 1994
A;Title: In vitro assembly of repertoires of antibody chains on the surface of phage by A;Reference number: 846390; MUID:94254092; PMID:8196048
A;Reference number: 846390
A;Reference preliminary
A;Reference preliminary
A;Residues: 1-114 <FIG>
A;Gross-references: UNIPARC:UPI00001137D6; EMBL:231688; NID:8499306; PIDN:CAA83493.1; PI
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Reywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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Best Local Similarity 82.5%; Pred. No. 3.6e-40;
Matches 104; Conservative 1; Mismatches 9; Indels 12; Gaps
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Protein Sequence Searches - February 2005

All of the sequence databases on ABSS have recently been updated.

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or members of the public who may encounter UniProt temporary accession numbers.

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GenCore version 5.1.6

Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: December 28, 2005, 13:44:24; Search time 159.923 Seconds

555.871 Willion cell updates/sec
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Title:

Perfect score: 668
Sequence: 1 QVQLVESCGGVVQPGRSLRL......NYYYYGMDVWGQGTTVTVSS 126
Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Gapop 10.0 , Gapext 0.5
Gapop 10.0 , Gapext 0.5
Searched: 2166443 Begs, 705528306 residues

Total number of hits satisfying chosen parameters: 2166443

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : UniProt 05.80:*

1: uniprot sprot:*

2: uniprot_trembl:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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O6MZV6_HUMAN O6GMW7_HUMAN	Q9HCC1 HUMAN HV3T HUMAN	QSM7V3_RAT O68CN4_HUMAN	O9UL72_HUMAN	Q5BK12_RAT	Q6MZV7 HUMAN	Q6P6C4_HUMAN	O6MZQ6 HUMAN	Q99KA4 MOUSE	Q96K68 HUMAN	Q6PDB8_MOUSE
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33	8 E	36	38	39	40	41	42	43	44	4.5

ALIGNMENTS

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MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
X Strausberg K.L., Felingold E.A., Groute L.H., Derge J.G.,
Attausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
Attausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
Attausner R.D., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A Diatchenko L., Marusina K., Parmer A.A., Rubin G.M., Hong L.,
A Diatchenko L., Marusina K., Perers G.J., Abramson R.D., Mullahy S.J.,
B Capleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunbaratne P.H.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunbaratne P.H.,
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Nilalon D.K., Muzny D.M., Sodergren E.J., Lu X., Glbbs R.A.,
A Wilting M., Madan A., Young A.C., Shevchenko Y., Boutfard G.G.,
Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,
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Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
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                                                                                                                                                                          Homo sapiens (Human).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Buarchontoglires, Primates, Catarrhini, Hominidae,
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Preferential utilization of conserved immunoglobulin heavy chain
variable gene segments during human fetal life.";
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NIH MGC Project;
Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases.
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PubMed=1383695; DOI=10.1016/0161-5890(92)90173-U;
Cuisinier A.M., Fumoux F., Fougereau M., Tonnelle C.;
                                                                        Created)
Last sequence update)
Last annotation update)
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                                   613 AA.
                                   PRT;
                                                                        01-MAR-2002 (TrEMBLrel. 20,
01-MAR-2002 (TrEMBLrel. 20,
01-MAR-2004 (TrEMBLrel. 26,
                                   QBWUK1_HUMAN PRELIMINARY;
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                                                                                                           Restricted utilization of germ-line VH3 genes and short diverse third
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81.2%; Pred. No. 2.2e-44;
ive 4; Mismatches 10; Indels
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PROSITE; PS50835; IG_LIKE; 5.
PROSITE; PS00290; IG_MHC; UNKNOWN_3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           to staphylococcal protein A.";
J. Exp. Med. 178:331-336(1993).
EMBL: BC02040; AAH20240.1; -; mRNA.
PIR; F36005; F36005.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PIR; G36005; G36005.

PIR; PH1642; PH1642.

PIR; PH1644; PH1644.

PIR; PH1646; PH1646.

PIR; PH1646; PH1646.

PIR; PL0120; PL0120.

PIR; S1119; S11116.

PIR; S31116; S31116.

PIR; S31116; S31116.

PIR; S31116; S31119.

PIR; S31116; S31119.

PIR; S3116; S31119.

PIR; S11119; S31119.

PIR; PL0161; IADQ.

SMR; Q8WUXI, 20-242.

SMR; Q8WUXI, 20-242.

SMR; Q8WUXI, 20-242.

INCEPPO; IPR003109; IG_MC.

INCEPPO; IPR003597; IG_MC.

INCEPPO; IPR003596; IG_WC.

INCEPPO; IPR003596; IG_WC.

INCEPPO; IPR003596; IG_WC.

INCEPPO; IPR003596; IG_WC.
                                                                      PubMed=1730252;
Raaphorst F.M., Timmers
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Best Local Similarity
Matches 104; Conserv
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                                                                                                Schuurman R.K.;
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Gaps

10;

Conservative

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61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARD--RGIAAGGNYYYYGMDVWGQ 118
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Mammalia, Butheria, Euarchontoglires, Primates, Catarrhini, Hominidae,
                           20 QVQLVESGGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYDGSNKYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            80 ADSVKGRPTISRDNGKNSLYLQMNSLRAEDTALYYCAKEIG---AHNFYYYGMDVWGQGT
1 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Osanger A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wambutt R., Heubner D., Mewes H.W., Weil B., Amid C., Os Fobo G., Han M., Wiemann S.; Submitted (JAN-2005) to the EMBL/GenBank/DDBJ databases. EMBL; BSK40627; CAE45781.1; -; mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS00230; IG_MHC; UNKNOWN_2.
Hypothetical protein.
SEQUENCE 472 AA; 51724 MW; 26CB340D0046D279 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                            05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein DKFZp686F15220.
Name=DKFZp686F15220;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 QUL031 HUMAN PRELIMINARY; PRT; 116 AA. 09UL03; 01-MAY-2000 (TrEMBLrel. 13, Created) 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                              472 AA
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Interpro; IPR007110; Ig-11ke.
Interpro; IPR007110; Ig c1.
Interpro; IPR003590; Ig MHC.
Interpro; IPR003596; Ig MHC.
Interpro; IPR003596; Ig WHC.
SMART; SM00409; IG; 2.
SMART; SM00407; IGc1; 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The German cDNA Consortium;
                                                                                                                                                                                                                                                                                                                                                                                              QEN089_HUMAN PRELIMINARY;
                                                                                                                                                                                                                                                            132 GTMVTVSS 139
                                                                                                                                                                                                        119 GTTVTVSS 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (Human)
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Q9UL93 HU
ID Q9UL
AC Q9UL
DT 01-M
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Name=scFv;
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                                                                               Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Buarchontoglires, Primates, Catarrhini, Hominidae,
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Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pubmed=2840480;

A Bird J., Galili N., Link M., Stites D., Sklar J.;

Bird J., Galili N., Link M., Stites D., Sklar J.;

T "Continuing rearrangement but absence of somatic hypermutation in immunoglobulin genes of human B cell precursor leukemia.";

J. Exp. Med. 168:229-246 [1988].

R PRIS, PH1644;

PIR; PL0120; PH0642.

R PSSP; P01772; 2F84.

SMR; Q9UL93; 1-116.

SMR; Q9UL93; 1-116.

R InterPro; IPR003196; Ig-like.
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                                                                                                                                                                                                                                                                                                                                                              NUCLEOTIDE SEQUENCE.

MEDILTNE-93301610; PubMed-8315388; DOI=10.1084/jem.178.1.331;
MILLSON J.L., Karr N.S., Oppliger I.R., Mannik M., Sasso E.H.;

"The structural basis of germline-encoded VH3 immunoglobulin binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                             "Myosin-reactive autoantibodies in rheumatic carditis and normal
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                                                                                                                                                                                    NUCLEOTIDE SEQUENCE.
MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531; MU X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M., Young D.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 116;
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10; Indels
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Last annotation update)
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81.6%; Pred. No. 2.9e-44;
iive 4; Mismatches 10
                                                                                                                                                                                                                                                                                                                       Clin. Immunol. Immunopathol. 87:184-192(1998).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                to staphylococcal protein A.";
J. Exp. Med. 178:331-336(1993)
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QGGMY2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches 102; Conservative
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05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
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                                                           Homo sapiens (Human).
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                                                                                                                                             NCBI_TaxID=9606;
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Name=IGHM;
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AC QG
AC QG
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TISSUB-Primary B-Cells;

WEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

A Straubberg R.L.; Feingold E.A.; Grouse L.H.; Derge J.G.;

A Lischul S.P.; Collins F.S.; Magner L., Schaefer C.P.; Bhat N.K.;

A Lischul S.P.; Zeeberg B., Buetow K.H.; Schaefer C.P.; Bhat N.K.;

A Distchenko L.; Marusina K., Farmer A.A.; Wang J.; Hafeh F.;

Diatchenko L.; Marusina K., Farmer A.A.; Rubin G.M.; Hong L.;

A Stapleton M.; Soares M.B.; Bonaldo M.F.; Casavant T.L.; Scheetz T.B.;

B Rownstein M.J.; Usdin T.B.; Toshiyuki S.; Carninci P.; Prange C.;

A Raha S.S.; Loquellano N.A.; Peters G.J.; Abramon R.D.; Mullahy S.J.;

B Bosak S.A.; McEran K.J.; Malek J.A.; Gunaratne P.H.;

Richards S.; Worley K.C.; Hale S.; Garcia A.M.; Gay L.J.; Hulyk S.W.;

A Nilalon D.K.; Muzny D.M.; Sodergren E.J.; Lu X.; Gibbs R.A.;

Rahesley J.; Helton E., Ketteman M.; Madan A.; Rodriques S.; Sanchez A.;

Whiting M.; Madan A.; Young A.C.; Shevchenko Y.; Bouffard G.G.;

B Blakesley R.W.; Touchman J.W.; Green E.D.; Dickson M.C.;

B Butterfield Y.S.N.; Krzywinski M.I.; Skalska U., Smailus D.E.;

R Generation and initial analysis of more than 15,000 full-length human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAR-DRGIAAGGN-----YYYYG 112
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76.1%; Pred. No. 2.2e-43;
ive 8; Mismatches 16; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE-Primary B-Cells;
NIH MGC Project;
Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.
EMBL; CONTATSS; AAH73758.1; -; mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROSITE; PS50835; IG LIKE; 5.
PROSITE; PS00290; IG WHC; UNKNOWN 3.
SEQUENCE 606 AA; 66185 MW; B6B38B51114E4C55 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
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Last annotation update)
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InterPro; IPR003599; Ig.
InterPro; IPR003599; Ig-1ike.
InterPro; IPR003597; Ig-1ike.
InterPro; IPR003596; Ig-MHC.
InterPro; IPR003596; Ig-MHC.
InterPro; IPR003596; Ig-W.
SMART; SM00409; IG-1; 4.
SMART; SM00400; IGC1; 4.
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Q65ZC9;
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Matches 102; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and mouse cDNA sequences.
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InterPro; IPR007110; Ig-like.
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Raaphorst R.M., Timmers E., Kenter M.J., Van Tol M.J., Vossen J.M.,
Schuurman R.K.;
Restricted utilization of germ-line VH3 genes and short diverse third
complementarity-determining regions (CDR3) in human fetal B lymphocyte
immunoglobulin heavy chain rearrangements.";
Bur. J. Immunol. 22:244-251(1992).
EMBL; AF035024; AAD56260.1; -; mRNA.
PIR; ST8486; ST8486.
HSSP; P01772; 2F848.
SMR; Q9UL90; 1-113.
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6
                                                                                                   STRAIN=CIG/7;

MEDLINE=9736-27799, PubMed=9219263; DOI=10.1038/nbt0797-629;

MEDLINE=9736-27799, PubMed=9219263; DOI=10.1038/nbt0797-629;

M Kontermann R.E., Wing M.G., Winter G.;

I "Complement recruitment using bispecific diabodies.";

I Nat. Biotechnol. 15:629-631(1997).

R EMBL; Y13056, CAA73499-11.; mRNA.

R InterPro; IPR003599, IG.

R InterPro; IPR003599, IG.

R InterPro; IPR003596, IG.

R InterPro; IPR003596, IG.

R SMART; SM00409; IG. 2.

R SMART; SM0406; IGV: 2.

R PR0SITE; PS50835; IG_LIKE; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 77.0%; Score 514.5; DB 2; Length 240; 80.2%; Pred. No. 1.7e-43; ive 6; Mismatches 10; Indels 9;
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MEDLINE-98277139; PubMed-9614934; DOI=10.1006/clin.1998.4531;
Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-WAY-2000 (TrEMBLrel. 13, Created)
WAY-2000 (TrEMBLrel. 13, Last sequence update)
01-WAY-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable region
                                                                                                                                                                                                                                                                                                                                                                                                     240 240 2569 MW; FDCFD3645F64B373 CRC64;
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Q9UL90;
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Best Local Similarity 80.2
Matches 101; Conservative
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                                       NCBI_TaxID=9606;
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MEDINES 2280825; PubMed=12477932; DOI=10.1073/pnas.242603899; MEDINES 2280825; PubMed=12477932; DOI=10.1073/pnas.242603899; MEDINES 2380825; PubMed=12477932; DOI=10.1073/pnas.242603899; MEDINES 2380825; PubMed=12477932; DOI=10.1073/pnas.242603899; MEDINES 2380825; PubMed=12477932; DOI=10.1073/pnas.242603899; MEDINES 2380825; MEDINE F.S., Zechaefer C.F., Bhat N.K., Altschul S.F., Zecheefer G.F., Marusina K., Farmer A.A., Rubin G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheefer T.B., Stapleton M.J., Uddin T.B., Peters G.J., Abramson R.D., Mullahy S.J., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Abramson R.D., Mullahy S.J., Abramson R.D., Mullahy S.J., Abramson R.D., Mullahon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A., Villaton D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A., Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., Abramson M.M., Roderson M.C., Shevchenko Y., Bouffard G.G., Abramson M.M., Raden B.D., Dickson M.C., Abramson M.M., Roderiquez A.C., Grimwood J., Schmutz J., Myers R.M., Schhein J.B., Jones S.J.M., Marra M.A., Schhein J.B., Jones B.D., Jones C.D., Jones L.D., Jones C.D., Jones
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Mammalia, Butheria, Buarchontoglires, Primates, Catarrhini, Hominidae,
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PubMed=1555592;
Makiya R., Stigbrand T.;
"Placental alkaline phosphatase has a binding site for the human
                                                                                                                                                                                                                                                                                                                                                                                                                         13;
                                                                                                                                                                                                                                                                                                                          76.9%; Score 513.5; DB 2; Length 113; 79.4%; Pred. No. 9e-44; ive 5; Mismatches 8; Indels 13;
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Director MGC Project;
Submitted (JAN-2002) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                         113 AA; 12437 MW; ED57FDD19086D07F CRC64;
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Last annotation update)
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Interpro; IPR003596; Ig.v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
NON TER 113 113
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01-MAR-2002 (TrEMBLrel. 20,
01-MAR-2004 (TrEMBLrel. 26,
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IPR003596; Ig_v.
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Matches 96; Conserv
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Mammalia; Butheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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Amino acid sequence of the variable region of a human mu chain:
location of a possible JH segment.";
Proc. Natl. Acad. Sci. U.S.A. 77:3239-3243(1980).
-!- MISCELIANBOOUS: This mu chain was isolated from the plasma of patient with macroglobulinemia.
-!- SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 573;
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                                                                                                                                                                                                                                                                                                                                                                                                                                76.3%; Score 510; DB 2; 76.6%; Pred. No. 1.3e-42; iive 12; Mismatches 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Homo bapiens (Human).
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SMR; P01766; 2-122.
GO; GO:0005576; C:extracellular region; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
                                                                                                            HSSP, P18529, 118K.

Ensembl; ENSG0000196122; Homo sapiens.
InterPro; IPR00110; Ig-like.
InterPro; IPR003597; Ig_c1.
InterPro; IPR003596; Ig_WC.
InterPro; IPR003596; Ig_W.
Pfam; PR0047; Cl-set; I.
Pfam; PR0047; Ig, 2.
SMART; SM00406; IGv; I.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS50835; IG_LIKE; 4.
immunoglobulin-G Rc portion.";
Bur. J. Biochem. 205.341-345(1992).
EMBL: BC021276; AAH21276.1; -; mRNA.
PIR; S21205; S31205.
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Matches 98; Conservative
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P01768;
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61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGNYYYYGMDVWGQGT 120
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cao J., Vescio R.A., Rettig M.B., Hong C.H., Kim A., Lee J.C.,
Lichtenstein A.K., Berenson J.R.;
"A CD10-positive subset of mailgnant cells is identified in multiple
myeloma using PCR with patient specific immunoglobulin gene primers.";
Leukemia 9:1948-1953(1995).
EMBL; S80866; AAD14339-1; -; mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                 1 QVELVESCGGVVZPGRSLRLSCAASGFTFSNYAMHWVRQPPGKGLEWVAVISYBGBBKYY
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G0; G0:005887; C:integral to plasma membrane; NAS.
G0; G0:0016066; P:cellular defense response (sensu Vertebrata); NAS.
InterPro; IPR001596; P:cellular defense response (sensu Vertebrata); NAS.
InterPro; IPR003596; Ig_v.
SMART; SMO0406; IGv.
NON_TER.
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                                                                                                                                         Pyrrolidone carboxylic acid.
                                                                                                                                                                122 122
122 AA; 13668 MW; A42D0F17D252F1C2 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             147 AA; 15768 MW; 8489FCAAA7BC925C CRC64;
                                                 Direct protein sequencing; Immunoglobulin domain; Immunoglobulin V region; Pyrrolidone carboxylic acid. DOMAIN 1 112 Ig-like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                  ch 75.9%; Score 507; DB 1; Similarity 76.2%; Pred. No. 4.5e-43; 96; Conservative 14; Mismatches 12
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MEDLINE=96071149; PubMed=7475288;
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
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STANDARD;
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126
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                       118 OGTTVTVSS 126
                                             140 KGTTVTVSS 148
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                                                                                                                                                                                                                           NCBI TaxID=9606;
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P01772;
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                                                                                                                   TISSUB-Primary B-Cells.

WEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

A Straubberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Rlausner R.D., Colling F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

B Rownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

B Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunarathe P.H.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

Rahey J., Helton E., Ketreman M., Madan A., Rodrigues S., Sanchez A.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

B Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

B Cheneration and initial analysis of more than 15,000 full-length human
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                                                                                                                                                             Homo sapiens (Human).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Buarchontoglires, Primates, Catarrhini, Hominidae,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (1952)

RMBL; BC041037; AAH41037.1; -; mRNA.

RMBL; BC041037; AAH41037.1; -; mRNA.

HSSP; PO1861; 1ADO.

InterPro; IPR003599; Ig.

InterPro; IPR003599; Ig.

InterPro; IPR003596; Ig.

InterPro; IPR003506; Ig.MHC.

InterPro; IPR003506; Ig.WHC.

InterPro; IPR003506; Ig.WHC.

InterPro; IPR003506; Ig.WHC.

InterPro; IPR003596; Ig.WHC.

InterPro; IPR003596; Ig.WHC.

InterPro; IPR003596; Ig.WHC.

InterPro; IRR003596; Ig.WHC.

InterPro; IRR003506; Ig.WHC.

InterPro; IRR003506; Ig.WHC.

InterPro; IRR003506; Ig.WHC.

InterPro; IRR00409; Ig.2.

SWART; SW00409; IGG1; 3.
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PROSITE; PS00290; IG MHC; UNKNOWN 2.
SEQUENCE 478 AA; 52667 MW; 17BED38D917970D6 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                   Last sequence update)
Last annotation update)
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                                                                               PRT;
                                                                                                      Created)
                                                                                                    05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27,
                                                                          QEPIB1_HUMAN PRELIMINARY;
QEPIB1;
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nes 98; Conservative
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NIH MGC Project;
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121 TVTVSS 126
                      121 LVTVSS 126
                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE
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                                                                                                                                                                                                            NCBI_TaxID=9606;
                                                                                                                                          IGHM protein.
Name=IGHM;
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This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
80 VDSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCAREFESTMTTVNADYYYFYMDVWG 139
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Eukaryote; Metazoa, Chordata; Craniata; Primates; Catarrhini; Hominidae;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Schmidt W.E., Jung H.-D., Palm W., Hilschmann N.; schmidt W.E., Jung H.-D., Palm W., Hilschmann N.; structure determination of antibodies. Primary structure of crystallized monoclonal immunoglobulin IgG1 KOL, I."; Hoppe-Seyler's Z. Physiol. Chem. 364:713-747(1983).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and 1.0-A resolution.";
J. Mol. Biol. 141:369-391(1980).
-!- SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain.
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PROSITE; PSSOSE; IG LIKE; 1.
3D-structure; Direct protein sequencing; Immunoglobulin domain;
Immunoglobulin V region; Pyrrolidone carboxylic acid.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Homo sapiens (Human).
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PDB; 2IG2; X-ray; H=2-126.
GO; GO:0005576; C:extracellular region; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
                                                                                                                                                                                                                                                                                                                                                                            126 AA
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MEDLINE=81072295; PubMed=7441755;
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MEDLINE=83289131; PubMed=6884994;
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61 ADSVKGRFTIFRDNSKNMMDLQMNSLRAEDTAVYYCAKDERGRLVGTYF----DYWGQGT 116
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Nucleic Acids Res. 17:4385-0(1989).
RMBL; BCOL5760; AAH15760.1; -; mRNA.
PIR; S05271; S05271.
PIR; S24260; S24260.
HSSP; P01861; 1AADQ.
Ensembl; ENSG00000130076; Homo sapiens.
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PROSITE, PS50835, IG_LIKE, 5.
PROSITE; PS00290; IG_MHC; UNKNOWN_3.
                                                                                                                                                                                                                                               PRT;
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InterPro; IPR003597; Ig cl.
InterPro; IPR003006; Ig-MHC.
InterPro; IPR003596; Ig-W.
Pfam; PF07654; Cl-Bet; 4.
                                                                                                                                                                                                                                            Q96BB9 HUMAN PRELIMINARY;
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NIH MGC Project;
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SEQUENCE 597 AA; 65
                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human).
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                                                        121 TVTVSS 126
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                                                                                                           117 LVTVSS 122
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                                                                                                                                                                                                                                                                                                                                                                                    protein.
                                                                                                                                                                                                                                                                                                                                                                                                                Name=IGHM;
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Mammalia; Butheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 QVQLVESGGGVVQPGRSLRLSCSSGGFIFSSYAMYWVRQAPGKGLEWVAIIWDDGSDQHY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 72.6%; Score 485; DB 2; Length 122; 77.0%; Pred. No. 7.3e-41; rive 6; Mismatches 19; Indels
                                                                                                                                                                                                                                                                                                                                 Length 126;
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01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable region
                                                                                                                                                                                                                                                                                                                              73.8%; Score 493; DB 1; Length 12
76.2%; Pred. No. 1.2e-41;
ive 11; Mismatches 19; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              122 AA; 13579 MW; 36054D41366545B8 CRC64;
                                                                                                                                                                                                                                                                           126 AA; 13718 MW; E4D71B52B16F8776 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cin. Immunol. Immunopathol. 87:184-192(1998).
EMBL; AP035030; AAD56266.1; -; mRNA.
HSSP; P01772; 2PB4.
SMR; O9UL64; 1-122.
Ensembl; ENSG0000130076; Homo sapiens.
InterPro; IPR00110; 19-1ike.
InterPro; IPR003596; 19-V.
SMART; SM00406; 1GV; 1.
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01-MAY-2000 (TrEMBLrel. 13, Last seq
01-OCT-2003 (TrEMBLrel. 25, Last ann
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                                                                                                                                                                                                                                                                                                                                                                                      96; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  121 TVTVSS 126
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TISSUB-Primary B-Cells;

XX TISSUB-Primary B-Cells;

XX Equaberg R.L., Feingold B.A., Groube L.H., Derge J.G.,

XX Equaberg R.L., Feingold B.A., Groube L.H., Derge J.G.,

XX Lausherg R.L., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

XX Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

XX Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

XX Expleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

XX Brownstein M.J., Usdin T.B., Toshiyuki S., Carninor P., Frange C.,

XX Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

XX Raha S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

XX Richards S.A., Walzuy D.M., Sodergren B.J., Lu X., Gibbs R.A.,

XX Richards S., Worlley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

XX Hands J., Helton B., Ketteman M.J., Madan A., Rodrigues S., Sanchez A.,

XX Mandan A., Young A.C., Shevchenko Y., Bouffard G.G.,

XX Blakesley R.W., Touchman J.W., Garen B.D., Dickson M.C.,

XX Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

XX Generation and initial analysis of more than 15,000 full-length human

XX and mouse CDNA sequences."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kishimoto T., Okajima H., Okumoto T., Taniguchi M.; "Nucleotide sequences of the CDNAs encoding the V-regions of H- and L-chains of a human monoclonal antibody with broad reactivity to
                                                                                                                                                                                                                                                                                                                                                                                                                Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65039 MW; 4FCA3AD8ECE263D9 CRC64;
                                                                                          01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
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Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002)
597 AA.
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HV3H_HUMAN
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MEDINE=229825; PubMed=12477932; DOI=10.1073/pnas.242603899;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
Altschul S.P., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
Altschul S.P., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
Altschul S.P., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
Altschul S.P., Jordan H., Moore T., Max S.I., Wang J., Haiteh F.,
A platchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A platchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A stapleton M., Scares M.B., Bonaldo M.F., Casavant T.L., Scheefer T.E.,
Brownstein M.J. Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Raha S.S., McZwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
Boakk S.A., McZwan P.J., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
A Halton D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
Mhiting M., Madan A., Young A.C., Shevchenko Y., Bouffaud G.G.,
A Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,
Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
B. Maria M.A., Schein J.B., Jones S.J.M., Marra M.A.,
                                                                                               2,
                                                                                                                                                                                                                                                                                                                                                                       61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARD-RGIAAGGNYYYYGMDVWGQG 119
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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    Length 597;
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EMBL; BC019046; AAH19046.1; -; mRNA.

HSSP; PO1861; 1ADQ.

SRR; Q6PJ95; DAPJ.

InterPro; IPR003599; Ig.

InterPro; IPR003599; Ig.

InterPro; IPR003506; Ig.Mc.

SMART; SMO0409; IG.1; 3.

SMART; SMO0409; IG.1; 3.

SMART; SMO0409; IG.1; 3.
                                                                                          97; Conservative 11; Mismatches 16; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002)
    DB 2;
72.5%; Score 484.5; DB 2 76.4%; Pred. No. 5.2e-40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and mouse cDNA sequences."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     QGPJ95_HUMAN PRELIMINARY;
QGPJ95;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rissum=Primary B-Cells;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TTVTVSS 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   138 TLVTVSS 144
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                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IGHG1 protein.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            120
    Query Match
                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HUMAN
                                                                                               Matches
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OGEOUSE HUD

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OC HOMO.

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This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
                                                                                                                                                                                                                                                                                                                           61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGNYYYYGMDVWGQGT 120
                                                                                                                                                                                                                                                                                                                                                       80 AASVKGRFTISRDNSKOTILSLQMNSLRVEDTAVYYCAKDQKPWYSNSWFLTNFDSWGRGT 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                               20 QAQLVESGGGGVVQPGSSLRLSCAASGFRFSNYGMHWVRQAPGKGLEWVAVFSYDESDKYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
                                                                                                                                                                 Gарв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  macroglobulin.
-!- SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain.
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0
                                                                                                         71.9%; Score 480; DB 2; Length 544; 74.6%; Pred. No. 1.3e-39;
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                                                                                                                                                              24; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HSSP; P01772; 2FB4.

SNR; P01775; 2FB4.

SNR; P01775; 5FB4.

GO; GO:000555; C:extracellular region; NAS.

GO; GO:0003823; F:antigen binding; NAS.

GO; GO:0005955; P:immune response; NAS.

InterPro; IFR007110; Ig-like.

InterPro; IFR007110; Ig-like.

INTERPRO; IRR018596; Ig-v.

SNART; SN00406; IGv; 1.

PROSITE; PSC0835; IG-LIKE; 1.

PROSITE; PSC0835; IG-LIKE; 1.

Inmunoglobulin V region; Pyrrolidone carboxylic acid.
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PROSITE; PS50835; IG LIKE; 4.
PROSITE; PS00290; IG MHC; UNKNOWN 2.
SEQUENCE 544 AA; 60102 MW; 1895814B2297C668 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      122 122
122 AA; 13167 MW; 74E5B6959E84100A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Homo sapiens (Human)
                                                                                                                                                              8; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=74175307; PubMed=4208843;
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Best Local Similarity 69.8%
Matches 88; Conservative
                                                                                                                                                                 94; Conservative
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Best Local Similarity 89.8
Matches 114; Conservative
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US-09-472-087-70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 QVQLVESGGGVVQPGRSLRL......NYYYYGMDVWGQGTTVTVSS 126
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/cgn2_6/ptodata/1/jaa/H_COMB.pep:*
/cgn2_6/ptodata/1/jaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/jaa/RE_COMB.pep:*
/cgn2_6/ptodata/1/jaa/RE_COMB.pep:*
                                          GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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US-09-848-798-134
US-09-424-840B-5
US-09-424-840B-16
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US-07-942-245-35
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                         protein search, using sw model
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seq length: 200000000
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US-09-240-274-26
US-09-240-274-26
Sequence 26, Application US/09240274
Patent No. 6255455
GENERAL INFORMATION:
APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
                                       14, Appl
89, Appl
139, Appl
139, App
77, Appl
77, Appl
77, Appl
79, Appl
79, Appl
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j Sequence 70, Application US/09472087

j Patent No. 6682736

j GENERAL INFORMATION:

APPLICANT: HANSON, DOUGLAS C.

APPLICANT: HANSON, DOUGLAS C.

APPLICANT: HANSON, BILLEN E.

APPLICANT: HANSON, STEVEN C.

APPLICANT: GILMAN, STEVEN C.

APPLICANT: GILMAN, STEVEN C.

APPLICANT: CORPORATION: CORPORATION: CORPORATION: CORPORATION: CORPORATION: USEOFFER C.

TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO CTLA-4

FILE REPERENCE: ABX-PPI

CURRENT APPLICATION NUMBER: US/09/472,087

CURRENT FILING DATE: 1999-12-23

PRIOR FILING DATE: 1999-12-23

NUMBER OF SEQ ID NOS: 147

SOFTWARE: PATENTIN VET. 2.1
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Pred. No. 8.1e-49;
1; Mismatches 9
US-09-472-087-80

US-08-862-124-17

US-08-862-124-14

US-09-269-332-89

US-09-240-274-139

US-09-840-459-77

US-09-840-459-79

US-09-497-625A-77

US-09-497-625A-79

US-09-315-926A-80

US-09-315-926A-80

US-09-240-274-16

US-09-848-798-145

US-09-848-798-145

US-09-848-798-145

US-09-848-798-145

US-09-848-798-145

US-09-848-798-145

US-09-848-798-16
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Sequence 153, Application US/09848798

| Patent No. 6858719
| GENERAL INFORMATION:
| APPLICANT: Siegel, Donald L. |
| TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF |
| TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF |
| TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF |
| TITLE OF INVENTION: BOSTING METHOD FOR PRODUCTION THEREOF |
| TITLE OF INVENTION: BOSTING METHOD FOR PRODUCTION THEREOF |
| TITLE OF INVENTION: BOSTING METHOD FOR PRODUCTION THEREOF |
| CURRENT FILING DATE: 2001-05-04 |
| PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/028,550 |
| PRIOR PRIOR APPLICATION NUMBER: EARLIER PHILING DATE: 1996-10-11 |
| NUMBER OF SEQ ID NOS: 224 |
| SOFTWARE: PATENTING DATE: 20 |
| SEQ ID NO 153 |
| LINGTH: 126 |
| TITLE OF INVENTION OF SEQ ID NOS: 224 |
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| TITLE OF INVENTION OF SEQ ID NOS: 224 |
| TITLE OF INVENTION OF SEQ
                                                                                                                                                                                                                                                                                                    APPLICANT: Siegel, Donald L.

TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: Rh(D)-BINDING PROPUCTION THEREOF
FILE REPERENCE: 09596-4202
CURRENT APPLICATION NUMBER: US/09/240,274
CURRENT PEPLICATION NUMBER: US/09/240,274
CURRENT PEPLICATION NUMBER: 60/081,380
EARLIER FILING DATE: 1998-04-10
EARLIER PILING DATE: 1996-04-10
EARLIER PILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
SOFTWARE: PATENTING DATE: 1996-10-11
LENGTHAR: 126
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83.7%; Score 559; DB 2; Length 126;
Best Local Similarity 83.3%; Pred. No. 4.1e-48;
Matches 105; Conservative 8; Mismatches 13; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 559; DB 2; Length 126; Pred. No. 4.1e-48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: anti-Rh(D) antibody clone SH56 US-09-240-274-153
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                                                                                                                                                                                                    Sequence 153, Application US/09240274 Patent No. 6255455 GENERAL INFORMATION:
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83.3%;
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ORGANISM: Homo sapiens
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Best Local Similarity
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Best Local Similarity
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GENERAL INFORMATION:
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
TILE REFERENCE: 09596-4202
CURRENT APPLICATION NUMBER: US/09/848,798
CURRENT FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/028,550
PRIOR APPLICATION NUMBER: EARLIER PILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
SOFTWARE: PALCHER PILING DATE: 2001-05-05
SEQ ID NO 26
LENGTH: 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGNYYYYGMDVWGQGT 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGNYYYYGMDVWGQGT 120
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   INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
                        FILE REFERENCE: 00596-4202
CURRENT APPLICATION NUMBER: US/09/240,274
CURRENT FILING DATE: 1999-01-29
FARLIER APPLICATION NUMBER: 60/081,380
EARLIER FILING DATE: 1998-04-10
EARLIER APPLICATION NUMBER: 60/028,550
FARLIER FILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 560
LENGTH: 126
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US-09-240-274-26
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; OTHER INFORMATION: anti-Rh(D) chain D31
US-09-848-798-26
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US-09-848-798-26
Sequence 26, Application US/09848798
; Patent No. 6858719
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ATTORNEY/AGENT INFORMATION:
NAME: Marscharg, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
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85.7%;
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Best Local Similarity 85.74
Matches 108; Conservative
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 25
LENGTH: 126
                                                                                                                                                                  Query Match
Best Local Similarity 83.33
Matches 105; Conservative
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                                                      TYPE: PRT
ORGANISM: Homo sapiens
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US-09-079-029-11
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APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2
CURRENT PEPLICATION NUMBER: US/9/848,798
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/240,274
PRIOR FILING DATE: EARLIER FILING DATE: 1999-01-29
PRIOR PILING DATE: EARLIER APPLICATION NUMBER: 60/028,550
PRIOR PILING DATE: EARLIER APPLICATION NUMBER: 60/028,550
                                                                                                                 61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGNYYYYGMDVWGQGT 120
                                                                                                                                    APPLICATE: Siegel, Donald L.

TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 05596-4207
CURRENT PELICATION NUMBER: 05/09/240,274
CURRENT PAPLICATION NUMBER: 05/081,380
EARLIER APPLICATION NUMBER: 60/081,380
EARLIER PILING DATE: 1999-04-10
EARLIER FILING DATE: 1999-04-10
EARLIER FILING DATE: 1996-10-11
SOFTWARE: PATENTION NUMBER: 60/028,550
EARLIER FILING DATE: 1996-10-11
SOFTWARE: PATENTION NUMBER: 05/028,550
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   13; Indels
   8; Mismatches
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US-09-240-274-25
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Matches 105; Conservative
     105; Conservative
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ORGANISM: Homo sapiens
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US-09-240-274-25
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LENGTH: 126
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Best Local
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Pred. No. 3.6e-47;
3; Mismatches 7; Indels
                                                          Score 557; DB 2; Length 126;
Pred. No. 6.4e-48;
8; Mismatches 13; Indel8
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CURRENY APPLICATION DATA:
APPLICATION NUMBER: US/09/079,029
; OTHER INFORMATION: anti-Rh(D) chain D30 US-09-848-798-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Adams, Camilia W.
APPLICANT: Ashkenazi, Avi J.
APPLICANT: Ashkenazi, Avi J.
APPLICANT: Churtharapai, Anan
APPLICANT: Kim, Kyung J.
TITLE OF INVENTION: Apo-2 Receptor
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: DNA Way
CITY: South San Francisco
STATE: California
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ORGANISM: Homo sapiens
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; Sequence 140, Application US/09240274
; Patent No. 6255455
; GENERAL INFORMATION:
; AFPLICANT: SIGGE1, Donald L.
; TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
; TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
; TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
; CURRENT APPLICATION NUMBER: US/09/240,274
; CURRENT FILING DATE: 1999-01-29
; EARLIER APPLICATION NUMBER: 60/081,380
; EARLIER FILING DATE: 1996-10-11
; NUMBER OF SEQ ID NOS: 224
; SOFTWARE: PatentIN Ver. 2.0
; SEQ ID NO 140
; LENGTH: 125
; TENGTH: 125
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Patent No. 685913

Patent No. 685914

Patent No. 685915

GENERAL INFORMATION:

APPLICANT: Siegel, Donald L.

TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

TURRENT FILING DATE: 2010-05-04

CURRENT APPLICATION NUMBER: US/09/848,798

CURRENT PILING DATE: 2010-05-04

PRIOR FILING DATE: EARLIER FILING DATE: 1999-01-29

PRIOR FILING DATE: EARLIER FILING DATE: 1996-10-11

NUMBER OF SEQ ID NOS: 224

SOFTWARE: PATENTIN OF SEC ID NOS: 224
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   40 QVQLVQSGGGVVQPGRSLRLSCAASGFIFSSYGMHWVRQAPGKGLEWVAGIFYDGGNKYY 99
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                                                                                                                                  100 ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARDRG
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US-09-848-798-140
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61 ADSVKGRFSISRDNSKNTLYLQVNSLRADDTAVYYCARDQR-AAAGIFYYSRMDVWGQGT 119
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                                                                                           Query Match 82.1%; Score 548.5; DB 2; Length 125; Best Local Similarity 83.3%; Pred. No. 4.5e-47; Matches 105; Conservative 9; Mismatches 11; Indels 1;
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; Patent No. 6790938
; GENERAL INFORMATION:
; APPLICANT: Bercht-Old, Peter
; APPLICANT: Escher, Robert F. A.;
; TITLE OF INVENTION: ANTI-GPIIB/IIIA RECOMBINANT ANTIBODIES
; FILE REFERENCE: 100564-0909
; CURRENT APPLICATION NUMBER: US/09/424,840B
; CURRENT FILING DATE: 1999-12-03
; PRIOR APPLICATION NUMBER: DE 19820663.1
; PRIOR PLILING DATE: 1998-05-08
; PRIOR APPLICATION NUMBER: DE 19755227.7
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Sequence 29, Application US/10330613A

Batent No. 6924360

GENERAL INFORMATION:
TITLE OF INVENTION: ANTIBODIES AGAINST THE MUC18 ANTIGEN
TITLE OF INVENTION: ANTIBODIES AGAINST THE MUC18 ANTIGEN
TITLE OF INVENTION: ANTIBODIES AGAINST THE MUC18 ANTIGEN
CURRENT APPLICATION NUMBER: 202-12-26
PRIOR APPLICATION NUMBER: 60/346299
PRIOR FILING DATE: 2001-12-18
NUMBER OF SEQ ID NOS: 90
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 29
; OTHER INFORMATION: anti-Rh(D) antibody clone SH16 US-09-848-798-140
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US-10-330-613A-29
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US-09-424-840B-6
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; ORGANISM: Homo sapiens
US-09-424-840B-16
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APPLICANT: Siggel, Donald L.

APPLICANT: Siggel, Donald L.

TITLE OF INVENTION: Rh (D) -BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

FILE REFERENCE: 09596-42U2

CURRENT PELLIG DATE: 1999-01-29

EARLIER APPLICATION NUMBER: 60/081,380

EARLIER PILING DATE: 1998-04-10

EARLIER PILING DATE: 1996-11

NUMBER OF SEQ ID NOS: 224

SOFTWARE PAPLICATION VERSE: 60/028,550

EARLIER FILING DATE: 1996-10-11

SOFTWARE PAPLICATION VERSE: 1996-10-11

SOFTWARE PAPLICATION NUMBER: 60/028,550

EARLIER FILING DATE: 1996-10-11

SOFTWARE PAPLICATION VERSE: 1996-10-11
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                                                                                                                                                                                                                     81.1%; Score 541.5; DB 2; Length : 84.1%; Pred. No. 2.2e-46; tive 6; Mismatches 11; Indels
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Best Local Similarity 82.7%; Pred. No. 4.9e-46;
Matches 105; Conservative 9; Mismatches 11; Indels
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US-09-240-274-13
PRIOR FILING DATE: 1997-12-12
PRIOR APPLICATION NUMBER: DE 19723904.8
PRIOR FILING DATE: 1997-06-06
NUMBER OF SEQ ID NOS: 128
SOFTWARE: Patentin version 3.1
SEQ ID NO 6
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US-09-848-798-13
; Sequence 13, Application US/09848798
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Patent No. 6255455
                                                                                                                                                                                                                     Query Match
Best Local Similarity 84.1<sup>3</sup>
Matches 106; Conservative
                                                                                                                                         TYPE: PRT
ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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118 TVTVSS 123
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US-09-424-840B-6
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US-09-240-274-13
                                                                                                                     LENGTH: 123
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APPLICANT: Siegel, Donald L.

ITILE OF INVENTION: RAI(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

FILE REFERENCE: 09596-4202
CURRENT PEPLICATION NUMBER: US/09/848,798

CURRENT PEPLICATION NUMBER: SA01-05-04

PRIOR PELICATION NUMBER: EARLIER APPLICATION NUMBER: 60/028,550

PRIOR FILING DATE: BARLIER FILING DATE: 1996-10-11

NUMBER OF SEQ ID NOS: 224

SOFTWARE: PATENTH OF THE PERIOR APPLICATION NUMBER: 60/028,550

SEQ ID NO 13

LENGTH: 126
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; Sequence 16, Application US/09424840B
; Sequence 16, Application US/09424840B
; GENERAL INFORMATION:
APPLICANT: Berchtchid, Peter
; APPLICANT: Berchtchid, Peter
; TITLE OF INVENTION: ANTI-GPIIB/IIIA RECOMBINANT ANTIBODIES
; TITLE OF INVENTION: ANTI-GPIIB/IIIA RECOMBINANT ANTIBODIES
; CURRENT APPLICATION NUMBER: US/09/424,840B
; CURRENT FILING DATE: 1999-12-03
; PRIOR FILING DATE: 1998-05-08
; PRIOR FILING DATE: 1991-12-12
; PRIOR FILING DATE: 1997-06-06
; NUMBER OF SEQ ID NOS: 128
; SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 80.5%; Score 538; DB 2; Length 126; Best Local Similarity 82.7%; Pred. No. 4.9e-46; Matches 105; Conservative 9; Mismatches 11; Indels
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Pred. No. 6.1e-46;
6; Mismatches 12;
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Best Local Similarity 82.7%;
Matches 105; Conservative
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120 TTVTVSS 126 ||||||| 118 TTVTVSS 124

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Search completed: December 28, 2005, 14:05:14 Job time : 36 secs

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TYPE: PRT
ORGANISM: Homo sapiens
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118 TVTVSS 123
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US-10-325-694-144
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US-10-292-088-117
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Sequence 14, App
Sequence 14, Appl
Sequence 14, Appl
Sequence 15, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 16, Appl
Sequence 116, Appl
Sequence 116, Appl
Sequence 116, Appl
Sequence 116, Appl
Sequence 25, Appl
Sequence 216, Appl
Sequence 118, Appl
Sequence 118, Appl
Sequence 121, Appl
Sequence 121, Appl
Sequence 121, Appl
Sequence 127, Appl
Sequence 127, Appl
Sequence 127, Appl
Sequence 117, Appl
Sequence 117, Appl
Sequence 117, Appl
Sequence 117, Appl
                                                                                                                                             ; Search time 119.538 Seconds (without alignments) 440.415 Million cell updates/sec
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                                                                                                                                                                                                                                                                          QVQLVESGGGVVQPGRSLRL.........NYYYYGMDVWGQGTTVTVSS 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Description
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Published Applications AA Main:*
(cgn2 6/ptodata/1/pubpaa/USOP_PUBCOMB.pep:*
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(cgn2 6/ptodata/1/pubpaa/USII_PUBCOMB.pep:*
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                   GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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US-10-325-694-144

US-10-364-743-14

US-10-364-743-98

US-10-452-593-14

US-10-452-593-14

US-10-371-942-82

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US-10-041-860-209

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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                December 28, 2005, 14:03:21
                                                                                                           - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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seq length: 200000000
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668
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No.
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254, App
1731, Ap
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17, App
17, App
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11, App
11, App
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23, App
23, App
31, App
31, App
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Publication No. US20030148463A1
GENERAL INFORMATION:
APPLICANT: KUFER, PETER
APPLICANT: RAUM, TOBIAS
TITLE OF INVENTION: NOVEL METHOD FOR THE PRODUCTION OF ANTI-HUMAN ANTIGEN
TITLE OF INVENTION: RECEPTORS AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
      Sequence 206, Sequence 218, Sequence 1731, Sequence 177, Sequence 17, Sequence 17, Sequence 245, Sequence 217, Seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 3
Sequence 1
Sequence 5
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US-10-727-155-206
US-10-727-155-218
US-00-727-155-218
US-00-880-748-1731
US-10-153-382-17
US-10-153-382-17
US-10-776-649-70
US-10-776-649-70
US-10-776-649-70
US-10-776-649-70
US-10-269-711-13
US-10-269-711-13
US-10-269-711-27
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US-10-269-711-27
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US-10-269-711-27
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US-10-269-711-31
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US-10-269-711-31
US-10-269-711-31
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APPLICANT: GLADUE, RONALD P.
APPLICANT: CGRVALAN, JOSE
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-HI
APPLICANT: FENG, XIAO
TITLE OF INVENTION: ANTIBODIES TO CD40
TITLE REFERRINCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/10/292,088
CURRENT FILING DATE: 2003-03-14
PRIOR PILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
SEQ ID NO 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-292-088-117
; Sequence 117, Application US/10292088
; Publication No. US20030211100A1
; GENERAL INFORMATION:
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Matches 114; Conservative
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FILE REFERENCE: 38164000

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                                                                                                                                                                                                                                   Score 584.5; DB 4; Length 138;
Pred. No. 3.5e-48;
4; Mismatches 10; Indels 1; Gaps
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86.7%; Pred. No. 4.8e-48;
tive 4; Mismatches 11; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION.
APPLICANT: Bowdish, Katherine S.
APPLICANT: Prederickson, Shana
APPLICANT: Prederickson, Shana
APPLICANT: Waruyama, Toshiaki
APPLICANT: Maruyama, Toshiaki
APPLICANT: Maruyama, Toshiaki
APPLICANT: No. 0520040009178Alan, Mary Jean
TITLE OF INVENTION: INMUNOTHERAPEUTICS FOR BIODEFENSE
FILE REFERENCE: 84 (1087-73)
CURRENT APPLICATION NUMBER: US 407484,743
CURRENT FILING DATE: 2003-02-11
PRIOR FILING DATE: 2002-11-25
NUMBER OF SEQ ID NOS: 118
SOPTWARE: Patentin version 3.2
SEQ ID NOS: 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              i LOCATION: (26)...(26)
i OTHER INFORMATION: xaa=unknown amino acid
US-10-364-743-14
CURRENT APPLICATION NUMBER: US/10/325,694
CURRENT FILING DATE: 2002-12-19
PRIOR APPLICATION NUMBER: US/09/403,107
PRIOR FILING DATE: 1999-10-14
NUMBER OF SEQ ID NOS: 152
SOFTWARE: Patentin Ver. 2.1
LENGTH: 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 14, Application US/10364743; Publication No. US20040009178A1
                                                                                                                                                                                                                                     87.5%;
                                                                                                                                                                                                                                 Query Match
Best Local Similarity 88.2
Matches 112; Conservative
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                                                                                                                                                         ; TYPE: PRT
; ORGANISM: HUMAN
US-10-325-694-144
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ORGANISM: human
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61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGN--YYYYGMDVWGQ 118
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Boadish, Katherine S.
APPLICANT: Wild, Martha A.
APPLICANT: Wild, Martha A.
APPLICANT: Mald, Martha A.
APPLICANT: Mald, Mary Jean
TITE OF INVENTION: IMMUNOTHERAPEUTICS FOR BIODEPENSE
FILE REPERENCE: 98 CIP (1087-73 CIP)
CURRENT APPLICATION NUMBER: US/10/452,593
CURRENT APPLICATION NUMBER: US 10/364,743
PRIOR FILING DATE: 2003-02-11
PRIOR FILING DATE: 2002-02-11
PRIOR FILING DATE: 2002-02-11
PRIOR FILING DATE: 2002-04-29
PRIOR PRILING DATE: 2002-04-29
PRIOR PRILING DATE: 2002-04-29
PRIOR PRILING DATE: 2002-11-25
NUMBER OF SEQ ID NOS: 118
SOFTWARE: PATENTIN VERSION 3.2
SEQ ID NO 14
LENGTH: 135
                                            GENERAL INFORMATION:
APPLICANT: Bowdish, Katherine S.
APPLICANT: Prederickson, Shana
APPLICANT: Wild, Martha A.
APPLICANT: Waruyama, Toshiaki
APPLICANT: Waruyama, Toshiaki
APPLICANT: Waruyama, Toshiaki
APPLICANT: Wo. US20040009178Alan, Mary Jean
TITLE OF INVENTION: IMMUNOTHERAPEUTICS FOR BIODEFENSE
FILE REFERENCE: 84 (1087-73)
CURRENT APPLICATION NUMBER: US/10/364,743
CURRENT APPLICATION NUMBER: US 60/428,807
PRIOR FILING DATE: 2002-11-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: MISC_FEATURE

1 DCCATION: (26)...(26)

1 OTHER INFORMATION: xaa=unknown amino acid

15-10-364-743-98
Sequence 98, Application US/10364743 Publication No. US20040009178A1
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                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 118
SOFTWARE: PatentIn version 3.2
SEQ ID NO 98
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|123 GTTVTVSS 130
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ORGANISM: human
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61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGN-YYYYGMDVWGQG 119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 150, Application US/10325694
Publication No. US20030148463A1
GENERAL INFORMATION
APPLICANT: KUFER, PETER
APPLICANT: RAUM, TOBIAS
TITLE OF INVENTION: NOVEL METHOD FOR THE PRODUCTION OF ANTI-HUMAN ANTIGEN
TITLE OF INVENTION: RECEPTORS AND USES THEREOF
FILE REPERENCE: 38164000
CURRENT APPLICATION NUMBER: US/10/325,694
CURRENT FILING DATE: 2002-12-19
PRIOR PILING DATE: 1999-10-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 QVQLVQSGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYDGSNKYY
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Pred. No. 5.5e-48;
3; Mismatches 9; Indels
                                                                          US-10-371-942-82

Sequence 82, Application US/10371942

Sequence 82, Application US/10371942

Publication No. US20030223994A1

GENERAL INFORMATION:

APPLICANT: Hoogenboom, Henricus Renerus Jacobus Mattheus

APPLICANT: Relter, Yoram

TITLE OF INVERTION: MHC-PEPTIDE COMPLEX BINDING LIGANDS

FILE REFERENCE: 10280-034001

CURRENT APPLICATION NUMBER: US/10/371,942

CURRENT PILING DATE: 2003-02-20

PRIOR FILING DATE: 2002-02-20
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Pred. No. 6.8e-48;
5; Mismatches 10
                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 121
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          87.1%;
88.9%;
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Best Local Similarity 87.4%;
Matches 111; Conservative
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Best Local Similarity 88.9
Matches 112; Conservative
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SOFTWARE: Patentin Ver. 2
SEQ ID NO 150
LENGTH: 138
                                                                                                                                                                                                                                                                                                                                                                                                                                               ; ORGANISM: Homo sapiens
US-10-371-942-82
123 GTTVTVSS 130
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Fublication No. US20040258699A1
GENERAL INFORMATION:
APPLICANT: Prederickson, Shana
APPLICANT: Prederickson, Shana
APPLICANT: Wild, Martha A.
CURRENT FILING DATE: 2003-06-02
PRIOR PLICATION NUMBER: US 10/364,743
PRIOR PLICATION NUMBER: US 60/356,086
PRIOR PLICATION NUMBER: US 60/376,408
PRIOR FILING DATE: 2002-04-29
PRIOR FILING DATE: 2002-04-29
PRIOR FILING DATE: 2002-04-29
PRIOR FILING DATE: 2002-04-29
PRIOR PLING DATE: 2002-04-29
PRIOR PLING DATE: 2002-11-25
NUMBER OF SEQ ID NOS: 118
SOFTWARE: PatentIn Version 3.2
                                                                                                                                                                Score 583; DB 5;
Pred. No. 4.8e-48;
4; Mismatches 11
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Pred. No. 4.8e-48;
4; Mismatches 11
                                     FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (26)...(26)
US-10-452-593-14
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86.7%;
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Best Local Similarity 86.7%;
Matches 111; Conservative
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TYPE: PRT
ORGANISM: human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: human
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Best Local 9
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Sequence 282, Application US/10041860; Publication No. US20030157109A1; GENERAL INFORMATION:
APPLICANT: Corvalan, Jose R.F.
APPLICANT: Jia, Xiao-Chi
APPLICANT: Fang, Xiao
APPLICANT: Fang, Xiao
APPLICANT: Chen, Francine
APPLICANT: Ghen, Francine
APPLICANT: Ghen, Francine
APPLICANT: Ghen, Richard
APPLICANT: Geati, Gadi
                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 209, Application US/10041860; Publication No. US20030157109A1; GENERAL INFORMATION:
Query Match 87.0%;
Best Local Similarity 88.1%;
Matches 111; Conservative 3
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US-10-041-860-209
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    61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKDMGWGSGWRPYYYYGMDVWGQG 120
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APPLICANT: OTALAILAN, JOSE R.F.
APPLICANT: Grad, Xiao-Chi
APPLICANT: Feng, Xiao-Chi
APPLICANT: Feng, Xiao-Dong
APPLICANT: Gazit, Gadi
APPLICANT: Gazit, Gadi
APPLICANT: Bezabeh, Binyam
APPLICANT: Bezabeh, Binyam
TITLE OF INVENTION: ANTIBODIES DIRECTED TO PDGFD AND USES
TITLE REFERENCE: ABGENIX.051A
CURRENT APPLICATION NUMBER: 2002-01-07
NUMBER OF SEQ ID NOS: 377
SOFTWARE: FastsEQ for Windows Version 4.0
SEQ ID NO 17
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                                                                                                                                                                      Sequence 1, Application US/10778915
; Publication No. US20050180979A1
; GENERAL INFORMATION:
; APPLICANT: DCCHER, MATTHIAS
; APPLICANT: PERMS, NADJA
APPLICANT: PRANG, NADJA
; APPLICANT: PRANG, NADJA
; TITLE OF INVENTION: Ant.-EpcaM Immunoglobulins
; TITLE REFERENCE: DEBE:031US
; CURRENT FILLING DATE: 2004-02-13
; NUMBER OF SEQ ID NOS: 2
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 1
; LENGTH: 457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Synthetic Peptide US-10-778-915-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17, Application US/10041860; Publication No. US20030157109A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
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; ORGANISM: homo sapiens
US-10-041-860-17
                                        TTVTVSS 126
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|121 TTVTVSS 127
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Best Local
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Score 581; DB 4; Length 126;
Pred. No. 6.9e-48;
3; Mismatches 12; Indels
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APPLICANT: Corvalan, Jose R.F.
APPLICANT: Jia, Xiao-Chi
APPLICANT: Feng, Xiao
APPLICANT: Feng, Xiao-Dong
APPLICANT: Chen, Francine
APPLICANT: Gazit, Gadi
APPLICANT: Bezabe, Binyam
TITLE OF INVENTION: ANTHEODIES DIRECTED TO PDGFD AND USES
TITLE OF INVENTION: ANTHEODIES
TOURENT PALLICATION NUMBER: US/10/041,860
CURRENT PALLICATION NUMBER: US/10/041,860
NUMBER OF SEQ ID NOS: 377
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 87.0%; Score 581; DB 4; Length 126; Best Local Similarity 88.1%; Pred. No. 6.9e-48; Matches 111; Conservative 3; Mismatches 12; Indels
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US-10-292-088-116
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Sequence 14, Application US/10665383

Sequence 14, Application US/10665383

Sequence 14, Application US/10665383

Sequence 14, Application US/10665383

APPLICANT: Floege, Juergen

APPLICANT: Gazit, Gadi

APPLICANT: LaRochelle, William

APPLICANT: Lichenstein, Henri

APPLICANT: Location Henri

APPLICANT: Lichenstein, Henri

PRIOR PILLING DATE: 2003-09-16

NUMBER OF SEQ ID NOS: 97

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 14

LENGTH: 126
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                                                                                                                                                                                                                                                                                                                 Length 126;
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TITLE OF INVENTION: ANTIBODIES DIRECTED TO PDGFD AND USES TITLE DEFENCE: ABGENIX.051A
CURRENT APPLICATION NUMBER: US/10/041,860
CURRENT FILING DATE: 2002-01-07
NUMBER OF SEQ ID NOS: 377
SSOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 282
LENGTH: 126
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Matches 111; Conservative
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CORGANISM: homo sapiens
US-10-041-860-282
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CORGANISM: homo sapiens
US-10-665-383-14
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61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGNYYYYGMDVWGQGT 120
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86.9%; Score 580.5; DB 4; Length 1:
Best Local Similarity 89.7%; Pred. No. 7.5e-48;
Matches 113; Conservative 1; Mismatches 9; Indels
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86.7%; Score 579; DB 4; Length 12.
Best Local Similarity 88.9%; Pred. No. 1.1e-47;
Matches 112; Conservative 2; Mismatches 10; Indels
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PUDLICATION NO. US20030211100A1

GENERAL INFORMATION:
APPLICANT: GLADUE, RONALD P.
APPLICANT: CORVALAN JOSE
APPLICANT: UIA, XIAO-CHI
APPLICANT: TERG, XIAO
TITLE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCES: AXE-PF/3 US
CURRENT APPLICATION NUMBER: US/10/292,088
CURRENT APPLICATION NUMBER: US/10/292,088
CURRENT APPLICATION NUMBER: 0/348,980
FRIOR FILING DATE: 2003-03-14
FRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: PATENTIN VOYER: 2.1
                                                                         APPLICANT: BEDILAN, VAHE
APPLICANT: GLADUE, RONALD P.
APPLICANT: GLADUE, JOSE
APPLICANT: GLADUE, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
TITLE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/10/292,088
CURRENT FILING DATE: 2003-03-14
PRIOR PPLICATION NUMBER: 60/348,980
PRIOR PLILING DATE: 2001-11-09
Sequence 116, Application US/10292088 Publication No. US20030211100A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 147
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; ORGANISM: Homo sapiens US-10-292-088-116
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CRGANISM: Homo sapiens
US-10-292-088-106
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WO200212502-A2
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Region
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Human 9E7
Human ant
Anti-inte
22B3 anti
22B3 anti
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Erythropo
Anti-PsaA
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Human IP1
16E1 anti
16E1 anti
Anti-inte
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Human mAb
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Aam51173 Human Vg/
Aau76334 Human ant
                                                  December 28, 2005, 13:43:04; Search time 126.462 Seconds (without alignments) 375.236 Million cell updates/sec
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                                                                                     US-09-920-137P-8
568
1 BIVLTQSPATLSLSPGERAT.....QQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                                                                        Description
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Adj73534
Adj73533
Adj73534
Adw07091
Adm51174
Adm51174
Adf11411
Adf11413
Adm41573
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GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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                                                                                                                                                     2443163 seqs, 439378781 residues
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                                                                                                                                                                                                                 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                  protein search, using sw model
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AAM51175
AAB72884
AEB01062
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ADF11435
ADM41547
ADB9877
ADS64653
ADM41573
ADM41573
ADF11395
ADF11395
ADF11419
ADF11419
ADF102886
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AAM51173
AAU76334
ADJ73533
ADJ73534
ADW07091
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genescqp2000s:*
genescqp2001s:*
genescqp2003s:*
genescqp2003as:*
genescqp2003as:*
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length: 2000000000
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                                                                                                 Perfect score:
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Maximum DB
                                                                                                          Sequence:
                                                                                                                                                      Searched:
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Result No.

Light cha IGF-IR an Anti-plat Human IP1 2D8 anti-

ADI22138 AEB01057 ADF11403

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Light cha

IgG kappa

Amino aci

Brythropo

Single ch

Brythropo

Anti-NGF-

SARS coro

Human IPI

Human IPI
                                                                                                                                                                                                                                                                                            Tumour necrosis factor alpha; TNF; antibody; light chain; CDR; complementarity determining region; antirheumatic; antiarthritic; antialicer; antiathmatic; antiallergic; antilnflammatory; antisickling; antidiabetic; antiartherosclerotic; vasotropic; antianginal; cardiant; antibacterial; virucide; fungicide; antileprotic; protozoacide; cytostatic; neuroprotective; antibacterial; virucide; fungicide; antileprotic; human; diagnosis; therapy.
                208
                                                                                                                                                                                                                                                                               Anti-tumour necrosis factor antibody light chain variable region.
Adf11415
Adf11439
Adf11439
Adf11439
Ab57362
Abr54916
Adb15292
Add15292
Add173532
Add73532
Add73532
Add73532
Add73532
Add73533
Add73532
Add73533
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                                                                                                                                                                                        ALIGNMENTS
ADF11415
ADF11439
ADF11439
ADF11439
ABF57362
ABF54916
ABF54916
ADF73528
ADF73528
ADF73532
                                                                                                                                                               AAB99371
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                                                                                                                                                                                                                                AAM51165 standard; protein; 108 AA
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/label= CDR2
57. .88
/label= FR3
89. .98
/label= CDR3
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29-SEP-2000; 2000US-0236826P.
01-AUG-2001; 2001US-00920137.
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/label= FR2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-AUG-2001; 2001WO-US024785
                                                                                                                                                                                                                                                                                                                                                                                                        4. .34
label= CDR1
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label= FR1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99. .108
/label= J6
                                                                                                                                                                                                                                                               (first entry)
(CENZ ) CENTOCOR INC
14-FEB-2002
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anti-tumour necrosis acator (TNR) antibody of the invention. The antibodies isolated human, primate, rodent, mammalian, chimeric, humanised and/or complementarity determining region (CDR)-grafted antiburantised and/or complementary used to the antibodies, immunoglobulins, cleavage products and other specified portions and variants, as well as anti-TNR antibody compositions, formulations, devices, transgenic animals, transgenic compositions, formulations, devices, transgenic animals, transgenic plants, and methods of making and using them. The anti-TNR antibody comprises at least a portion of an immunoglobulin molecule, especially the heavy chain and/or light chain variable regions given in AMMS1164 and in the present sequence, or either all of the CDRs of the heavy chain (see AAMS11858-60) or all of the CDRs of the light chain (see AAMS1161-53). The antibody may inhibit TNP-induced cell adhesion molecules, inhibit TNP binding to receptor, or provide Arthritic Index improvement in a mouse model. It is useful for diagnosing or treating a TNP related condition in a cell, tissue, organ or animal (claimed) such as rheumatoid arthritis, gastric ulcer, asthma, allergic rhinitis, Crohn's pathology, sickle cell anaemia, diabetes, a cardiovascular disease such as antiponed cardial infections, pneumonia, leprosy and malaria, a malignant viral, and fungal infections, pneumonia, leprosy and malaria, a malignant callengs. Parkinson's disease, spinal ataxia, Alzheimer's disease and creutzfeldt-Jakob disease, spinal ataxia, Alzheimer's disease and creutzfeldt-Jakob disease.
                                                                                                                                  isolated mammalian anti-tumor necrosis factor antibody, useful for
                                                                                                                                                                                                                                                                                                 The present sequence is that of the light chain variable region of an anti-tumour necrosis factor (TNF) antibody of the invention. The
                                                                                                                                                           treating sickle cell anemia, diabetes, atherosclerosis, restenosis, angina pectoris, myocardial infarction, leprosy.
                         Shealy D;
                         Scallon B,
                            Heavner G,
                                                                                                                                                                                                                                              Claim 9; Page 129-130; 131pp; English
                            Giles-Komar J, Knight DM,
                                                                                                                                     Novel
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98.4%; Score 559; DB 5; Length 108 99.1%; Pred. No. 1.1e-37; ive 0; Mismatches 1; Indels Matches 107; Conservative Local Similarity Sequence 108 AA; Query Match

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Gaps

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Length 108;

61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108 -

EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA 60

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RESULT 2

AAM51173 standard; protein; 108 AA

AAM51173;

(first entry) 10-JUN-2002

Human Vg/38K-type light chain variable region.

Tumour necrosis factor alpha; TNF; antibody; light chain; CDR; complementarity determining region; antirheumatic; antiarthritic; antialcer; antialcer; antialcer; antialcer; antialcer; antialcer; antidiaectic; antiartherosclerotic; vasorropic; antiangental; cardiant; antibacterial; virucide; fungicide; antileprotic; protozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosis; therapy; Vg/38K.

Homo sapiens

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Cregion encoded by a human germine gene in a transgenic mouse used in human monoclonal antibody (mAb) construction. A GenTMV fusion was performed using spleen cells from a hybrid mouse containing human corrected by a human germine gene in a transgens that was memonic burners and constant region antibody transgenes that was immunised with performed using spleen cells from a hybrid mouse containing human corrected by a single antibody transgenes that was immunised with recombinant human tumoru necrosis factor (TRV) alpha Human mabs were obtained that bound immobilised human TNF alpha with apparently high cotained that bound immobilised human TNF alpha with apparently high avidity. These mAbs had a totally human 1gd1, kappa isotype. The mature correct for the light chain variable region of 2 of the mAbs. TNV14 and portion of the light chain variable region of 2 others, TNV148(B) and TNV196 (see TNV15, were identical (see AAM51174) to the present sequence, while the complementarity determining region (CDR)-grafted anti-TNF antibodies, and cleavage products and variants, as well as anti-TNF antibody compositions, encoding or complementary uncleic acids, vectors, immunoglobulins, and cleavage productes and variants, as well as anti-TNF antibody compositions, of complementary uncleic acids, vectors, cantibody compositions, formulations, devices, transgenic animals, cransgenic plants, and methods of making and using them. The anti-TNF ransgenic plants, and methods of making and using them. The antibodies at least a portion of an immunoglobulin molecule, capped to the present sequence and in AAM51165, or either all of the CDRs of the CDRs of the present sequence and in AAM51165, or either all of the CDRs of the CDRs of the present sequence and in AAM51165, or either all of the CDRs of the CDRs of the present sequence and in AAM51165, or either all of the CDRs of the CDRs of the present in a mouse model. They are useful for diagnosing or treating a rheumatoid arthritis, gastric ulcer, asthm a llergic rhinity or any parth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel isolated mammalian anti-tumor necrosis factor antibody, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence is that of a human Vg/38K-type light chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                treating sickle cell anemia, diabetes, atherosclerosis, restenosis, angina pectoris, myocardial infarction, leprosy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shealy D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Scallon B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Giles-Komar J, Knight DM, Heavner G,
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Fig 5; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                         29-SEP-2000; 2000US-0236826P.
01-AUG-2001; 2001US-00920137.
                                                                                                                                                                                                                                                                                                                                                                                     07-AUG-2001; 2001WO-US024785
                                                                                                                                                                                                                                                                                                                                                                                                                       07-AUG-2000; 2000US-0223360P
                                                                                                                                                     CDR2
                                                                                                                                                                    57. .88
/label= FR3
                                                                                                                                                                                                            89. .98
/label= CDR3
                                                                         /label= CDR1
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                                                                                                               FR2
                                                                                                                                                                                                                                                99. .108
/label= J3
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/label= 1
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/label= (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CENZ ) CENTOCOR INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-217194/27.
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 Key
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel isolated mammalian anti-dual integrin antibody, useful for diagnosing or treating dual integrin related condition such as rheumatoid arthritis, gastric ulcer, asthma, atherosclerosis, restenosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; antibody; dual integrin; HC CDR; variable region; LC CDR; medical device; immune related disease; rheumatoid arthritis; gastric ulcer; asthma; allergic rhinitis; Crohn's pathology; sickle cell anaemia; diabetes; cardiovascular disease; arteriosclerosis;
                                                                                                                                                                                                                                                                HILLI SPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                                          1 BIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          atherosclerosis; restenosis; angina pectoris; myocardial infarction; infectious disease; pneumonia; leprosy; malaria; malignant disease; leukaemia; chronic myelocytic leukaemia; multiple myeloma; meurological disease; multiple sclerosis; Parkinson's disease; Alzheimer's disease; creutzfeldt-Jakob disease.
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                                                                                                                                                                                                                                                                                                                                      61 RPSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                         RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human anti-dual integrin antibody complete variable region #2.
                                                                                                              Length 108
                                                                                                                                                                   1; Indels
Alzheimer's disease and Creutzfeldt-Jakob disease
                                                                                                              Score 559; DB 5;
Pred. No. 1.1e-37;
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                                                                                          99.1%; Pred. No. ....
99.1%; Pred. No. ....
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAU76334 standard; peptide; 108 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-AUG-2001; 2001WO-US024784.
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01-AUG-2001; 2001US-00920267.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                   Matches 107; Conservative
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                                                                                                                                        Similarity
                                                       Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Giles-Komar J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200212501-A2
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                                                                                                              Query Match
                                                                                                                                              Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mimetic; CDR mimetibody; gene therapy; transgenic; immune;
cardiovascular; infectious; malignant; neurologic disease; anaemia;
immunomodulator; cardiant; antimicrobial; cytostatic; neuroprotective;
                                     such as bacterial,
                                                                                                                                                                                                                                                                                                                                                                       1 EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                                                                                                                                                                 1 BIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
               arteriosclerosis, atherosclerosis, restenosis, angina pectoris, my coardial infarction, infectious disease in a cell such as bacterial disease und fungal infections, pneumonia, leprosy, malaria; malignant disease such as leukaemia, chronic myelocytic leukaemia, Burkitt's lymphoma, multiple myeloma; neurological disease such as multiple sclerosis, Parkinson's disease, spinal ataxia, Alzheimer's disease, creutzfeldt-Jakob disease and many other diseases given in the specification. The present sequence is an anti-dual integrin human (AAU76327-AAU76332)
                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to novel mammalian CDR mimetibodies, specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           variable region comprising human framework or ligand binding region, useful for preparing a composition for treating e.g., immune, cardiovascular or neurologic disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New CDR mimetibody comprising a portion of a heavy or light chain
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sickle cell anaemia, diabetes, cardiovascular disease such as
                                                                                                                                                                                                                                                                                                                                                                                                                           61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                   Score 559; DB 5; Length 108;
Pred. No. 1.1e-37;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Erythropoietin light chain mimetibody SegID 989.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; SEQ ID NO 989; 97pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADJ73533 standard; protein; 108
                                                                                                                                                                                                                                                               98.4%;
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                                                                                                                                                                                                                                                                                                           Matches 107; Conservative
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                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                               Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003084477-A2.
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                                                              diseases, as well as anaemia. Accordingly, they exhibit immunomodulator, cardiant, antimicrobial, cytostatic and neuroprotective activities. This polypeptide sequence is an erythropoietin light chain mimetibody of the
that can be useful in gene therapy and the generation of transgenic plants and animals. Furthermore, the CDR mimetibody is useful for preparing compositions for modulating, treating or reducing the symptoms of immune, cardiovascular, infectious, malignant and/ or neurologic
                                                                                                                                                                                                                                                                   1 EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mimetic; CDR mimetibody; gene therapy; transgenic; immune;
cardiovascular; infectious; malignant; neurologic disease; anaemia;
immunomodulator; cardiant; antimicrobial; cytostatic; neuroprotective;
                                                                                                                                                                                                                                       BIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to novel mammalian CDR mimetibodies, specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New CDR mimetibody comprising a portion of a heavy or light chain variable region comprising human framework or ligard binding region, useful for preparing a composition for treating e.g., immune, cardiovascular or neurologic disease.
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                                                                                                                                                                           Length 108;
                                                                                                                                                                                                        1; Indels
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                                                                                                                                                                         98.4%; Score 559; DB 7;
99.1%; Pred. No. 1.1e-37;
ive 0; Mismatches 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                ADJ73534 standard; protein; 108 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                          Similarity
                                                                                                                                            Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003084477-A2
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                                                                                                                  invention.
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                                                                                                                                                                         Query Match
Best Local 8
                                                                                                                                                                                                          Matches
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portions or variants thereof. Specifically, it refers to an antibody fragment where a protein has been inserted into, or replaces a portion of or more CDR regions, such that each CDR mimetibody comprises at least one portion of a heavy chain framework region and at least one ligand binding region (LBR). The present invention describes human mimetibodies, including modified immunoglobulins and cleavage products that can be useful in gene therapy and the generation of transgenic plants and animals. Furthermore, the CDR mimetibody is useful for preparing compositions for modulating, treating or reducing the symptoms

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The invention describes an isolated binding member comprising at least one binding domain capable of specifically binding Streptococcus presentations aurface adhesin A (PsaA) protein, the binding domain having a dissociation constant Kd for PsaA which is less than 1 x10 -6 M. Also described are: an isolated mucleic acid molecule encoding at least a part of the binding member described above; a vector comprising the nucleic acid molecule of (1); a host cell comprising the nucleic of (1); a cell line engineered to express the binding member described above; detecting or diagnosing a disease or disorder associated with phenumococcus in an individual; a kit comprising at least one binding member described above, the antibody being labeled; and a pharmaceutical composition comprising at least one binding member is useful for producing a pharmaceutical composition for the treatment of Pneumococcus infection. The binding member, methods and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antibacterial; antiinflammatory; immunosuppressive; antibody engineering; pharmaceutical; infection; pneumonia; meningitis; sepsis; pneumococcal surface adhesin A; PsaA; light chain variable region.
                                                                                                                                                                                                                         9
of immune, cardiovascular, infectious, malignant and/ or neurologic diseases, as well as anemia. Accordingly, they exhibit immunomodulator, cardiant, antimicrobial, cytostatic and neuroprotective activities. This polypeptide sequence is an erythropoietin light chain mimetibody of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New binding member towards Streptococcus pneumoniae surface adhesin A protein, useful for treating or preventing pneumonia, meningitis and/or
                                                                                                                                                                                                                                              1 EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
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                                                                                                                                                                                                                                                                                                                       RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
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                                                                                                                                                 Score 559; DB 7; Length 108;
Pred. No. 1.1e-37;
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                                                                                                                                                                                       0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; SEQ ID NO 40; 137pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                ADW07091 standard; protein; 108
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                                                                                                                                                   98.4%;
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11-JUL-2003; 2003US-0486647P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-APR-2005 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anti-PsaA-antibody 9A7 VK
                                                                                                                                                   Query Match
Best Local Similarity 99.1
Matches 107; Conservative
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                                                                                                                Sequence 108 AA;
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                                                                            invention.
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Shealy

Scallon B,

Heavner G,

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Giles-Komar J, Knight DM,
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composition are useful for treating or preventing an individual suffering
from disorders or diseases associated with S. pneumoniae, e.g. pneumonia,
meningitis and/or sepsis. This is the amino acid sequence of anti-
pneumococcal surface adhesin A (PsaA)-antibody 9A7 kappa light chain
                                                                                                                  9
                                                                                                                                   9
                                                                                                                                                                                                                                                                                                     Tumour necrosis factor alpha; TNF; antibody; heavy chain; CDR; complementarity determining region; antitheumatic; antiarthritic; antiantiathratic; antiantializer; antializer; antializer; antializer; antializer; antiatherosclerotic; antiatherosclerotic; antiatherosclerotic; antiatherosclerotic; antiantial; cardiant; antibacterial; virucide; fungicide; antileprotic; protozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosis; therapy; TNV14; TNV15; monoclonal antibody; mab.
                                                                                                                               BIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                  1 BIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                   Human recombinant mAb TNV14, TNV15 light chain variable region.
                                                                                                                                                        RPSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                   RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFFFGPGTKVDIK 108
                                                                             Length 108;
                                                                           98.4%; Score 559; DB 9; Length 10 99.1%; Pred. No. 1.1e-37; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                .. .20
/label= Signal_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                   20. .129
/label= Mature protein
                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                           AAM51174 standard; protein; 129 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   109. .118
/label= CDR3
119. .129
/label= J3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-AUG-2000; 2000US-023360P.
29-SEP-2000; 2000US-0236826P.
01-AUG-2001; 2001US-00920137.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          55. .69
/label= FR2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               70. .76
/label= CDR2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-AUG-2001; 2001WO-US024785.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        44. .54
/label= CDR1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 77. .108
/label= FR3
                                                                                                                                                                                                                                                                                                                                                                                                                                                      21. .43
/label= FR1
                                                                                                                                                                                                                                                                 (first entry)
                                                                                                Matches 107; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CENZ ) CENTOCOR INC.
                                                                                       Local Similarity
                                        variable region.
                                                          Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200212502-A2
                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                  10-JUN-2002
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                                                                                                                                                                                                                                              AAM51174;
                                                                                                                                                         61
                                                                                                                                                                           61
                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                               Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                   Protein
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Compositions, factor (TMF) human recombinant monoclonal antibodies (mAbs) TWN14 and TWN15. These are human mabs produced from a GenTWV than on TWN15. These are human mabs produced from a GenTWV than on TWN15. These are human mabs produced from a GenTWV and constant region antibody transgenes that was immunised with human TWF alpha with high avidity alpha. The human mabs bind immobilised human TWF alpha with high avidity and have a totally human 1gG1, kappa isotype. The mature portion of the can have a totally human 1gG1, kappa isotype. The mature portion of the and germline sequence (see AAM51175) show a single amino acid difference. If it invention provides human, primate, mammalian, rodent, chimeric, chimanised and/or complementary nucleic acids, vectors, host calls.

The invention provides human, primate, mammalian, rodent, chimeric, thumanised and/or complementary nucleic acids, vectors, host calls.

TR antibodies, immunoglobulins, cleavage products and other specified portions and variants, as well as anti-TWF antibody compositions, formulations, dereshing them. The anti-try antibody compositions, formulations, dereshing and using them. The anti-TWF antibody compositions, formulation of an immunoglobulin molecule, especified plants, and methods of making and using them. The anti-TWF antibody compositions, formulation of an immunoglobulin molecule, especified condition in a call, tissue, organ or animal (claimed) such as created condition in a cell, tissue, organ or animal (claimed) such as the antibodies may inhibit TWF induced cell adhesion molecules, inhibit the mouse model. They are useful for diagnosing or treating a TWF copy sickle cell anaemia diabetes, a cardiovascular disease such as arteriosclerosis, atherosclerosis, restenosis, allergic rhinities, gastric ulcer, asthma, allergic rhinities, copy arcticular, and fungal infactions, permonia, leprosy and malaria, and multiple mycloma, or a neurological disease such as multiple mycloma, or a neurological disease such as and multiple mycloma, or a neur
                                                                                                                    Novel isolated mammalian anti-tumor necrosis factor antibody, useful for treating sickle cell anemia, diabetes, atherosclerosis, restenosis, angina pectoris, myocardial infarction, leprosy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9
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                                                                                                                                                                                                                                                                                                                                                                     present sequence is that of the light chain variable region of anti-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 129;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 559; DB 5; Length 12
Pred. No. 1.3e-37;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human mAb TNV148(B), TNV196 light chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAM51175 standard; protein; 129 AA
                                                                                                                                                                                                                                                                                   Example 3; Fig 5; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98.4%;
99.1%;
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WPI; 2002-217194/27.
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                                           N-PSDB; ABL53514.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-JUN-2002
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Matches 107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -
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Novel isolated mammalian anti-tumor necrosis factor antibody, useful for treating sickle cell anemia, diabetes, atherosclerosis, restenosis, angina pectoris, myocardial infarction, leprosy.
       antidiabetic; antiarteriosclerotic; antiatherosclerotic; vasotropic; antiatherosclerotic; vasotropic; antiatherosclerial; virucide; fungicide; antileprotic; protozcacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosis; therapy; TNV148(B); TNV196; monoclonal antibody; mAb.
antiulcer; antiasthmatic; antiallergic; antiinflammatory; antisickling;
                                                                                                                                                                                                                                                                                                                                                                             Shealy D;
                                                                                                                                                                                                                                                                                                                                                                             Scallon B,
                                                                                                                                                                                                                                                                                                                                                                             Heavner G,
                                                                                   . .20
Tabel= Signal_peptide
                                                                                                    20. .129
/label= Mature_protein
                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; Fig 5; 131pp; English.
                                                                                                                                                                                              77. .108
/label= FR3
109. .118
/label= CDR3
119. .129
/label= J3
                                                                                                                                                                                                                                                                                                                      07-AUG-2000; 2000US-0223360P.
29-SEP-2000; 2000US-0236826P.
01-AUG-2001; 2001US-00920137.
                                                                                                                                                                                                                                                                                                    07-AUG-2001; 2001WO-US024785
                                                                                                                                          44. .54
/label= CDR1
                                                                                                                                                                             70. .76
/label= CDR2
                                                                                                                                                          55. .69
/label= FR2
                                                                                                                       21. .43
/label= FR1
                                                                                                                                                                                                                                                                                                                                                                             Giles-Komar J, Knight DM,
                                                                                                                                                                                                                                                                                                                                                          (CENZ ) CENTOCOR INC.
                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-217194/27.
N-PSDB; ABL53515.
                                                                                                                                                                                                                                                                WO200212502-A2.
                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                   14-FEB-2002
                                                                                 Peptide
                                                                                                    Protein
                                                                                                                       Region
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tumour necrosis factor (TMF) human recombinant monoclonal antibodies
(mAbs) TNV148(B) and TNV196. These are human mabs produced from a GenTNV
fusion using spleen cells from a hybrid mouse containing human variable
and constant region antibody transgenes that was immunised with human TNF
alpha. The human mabs bind immobilised human TNF alpha with high avidity
and have a totally human IgG1, kappa isotype. The mature portion of the
cand have a totally human IgG1, kappa isotype. The mature portion of the
light chain variable region differs from that of the Vg/38-type light
chain germline sequence (see AAM51173) by a single residue (Tyr-50
replacing Ser). The light chain variable regions of 2 other human mabs
(see AAM51174) were identical to the germline sequence. The invention
provides isolated human, primate, rodent, mammalian, chimeric, humanised
and/or complementarity determining region (CDR)-grafted anti-TNF
andlors immunoglobulins, cleavage products and other specified
portions and variants, as well as anti-TNF antibody compositions,
encoding or complementary nucleic acids, vectors, host cells, compositions, formulations, devices, transgenic animals, transgenic plants, and methods of making and using them. The anti-TNF antibody comprises at least a portion of an immunoglobulin molecule, especially the heavy chain and/or light chain variable regions given in the present sequence and in AAM51165, or either all of the CDRs of the heavy chain (see AAM51158-60) or all of the CDRs of the light chain (see AAM51161present sequence is that of the light chain variable region of anti-The

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inhibit TNF binding to receptor, or provide Arthritic Index improvement in a mouse model. They are useful for disagnesing or treating a TNF related condition in a cell, issue, organ or animal (claimed) such as rheumatoid arthritis, gastric ulcer, asthma, allergic rhinitis, Crohn's pathology, sickle cell annemia, diabetes, a cardiovascular disease such as arteriosclerosis, atherosclerosis, restenosis, angina pectoris or myocardial infarction, an infectious disease in a cell such as bacterial, viral, and fungal infectious, pneumonia, leprosy and malaria, a malignant disease such as leukaemia, chronic myelocytic leukaemia, Burkitt's a malignant lymphoma and multiple myeloma, or a neurological disease such as multiple spelvaemia, parkinson's disease, spinal ataxia, Alzheimer's disease and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention provides the protein and coding sequences for human monoclonal antibodies which bind specifically to the HER2/neu growth factor receptor (also known as erbB2). These are designated 3-F2, 1-D2 and 2-B8. They can be used in the immunotherapy-based treatment and prognosis of cancers, particularly adenocarcinomas such as salivary gland, stomach, kidney, mammary gland, lung and squamous cell carcinomas, and ovarian cancer. The present sequence is part of an antibody of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; HER2; neu; erbB2; oncogene; cancer; antibody; immunotherapy; 3-F2;
                                                                                                                                                                                                                                                                                                                                                                                                                                              9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New human monoclonal antibody that specifically binds to growth factor receptor HER2/neu, for treating, preventing or diagnosing diseases characterized by aberrant HER2/neu expression e.g. cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                                                                                                                                                                                                                                                              1 BIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       81 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGFGTKVDIK 128
                                                                                                                                                                                                                                                                                                                                                         Length 129;
                                                                                                                                                                                                                                                                                                                                                       98.4%; Score 559; DB 5; Length 12
99.1%; Pred. No. 1.3e-37;
ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human anti-HER2/neu antibody 2-E8 light chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 110; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1-D2; 2-E8; growth factor receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB72884 standard; protein; 108 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JUL-1999; 99US-0146313P.
10-MAR-2000; 2000US-0188539P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                    Creutzfeldt-Jakob disease
                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 99.1
hes 107; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-168698/17.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB; AAF75590
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                                                                                                                                                                                                                                                                                                                Sequence 129 AA;
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                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                    Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 9
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Homo
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88888888888888888888888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated human monoclonal antibody or an antigen-binding portion, which specifically binds to human interferon gamma inducible protein 10 (IP-10), useful for treating viral or bacterial infection, or inflammatory or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to an isolated human monoclonal antibody or an antigen-binding portion, which specifically binds to human IP-10 and exhibits at least one property selected from: inhibits binding of IP-10 to CKR3; inhibits IP-10 induced calcium flux; inhibits IP-10 induced cross-react with human ITAC. The isolated human monoclonal antibody is useful for treating an inflammatory or autoimmune disease including multiple sclerosis, rheumatoid arthritis, inflammatory bowel disease erythematoeus, Type I diabetes, inflammatory skin disorders (e.g., psoriasis, lichen planus), autoimmune thyroid disease (e.g., Graves'
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                                                                                                                                                                                                                                                                                                                                                               antibody; IP-10; inflammation; immune disorder; dermatological disease;
                                                                                                                        EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPQQAPRLLIYDASNRATGIPA
                                                                                                       EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                                                                                                                                                                                         respiratory disease, neurological disease, degeneration, infection, Neuroprotective, Antiarthritic, Antinheumatic, Antinflammatory; Gastrointestinal-Gen.; Antiulcer, Dermatological, Immunosuppressive, Antidiabetic, Antipsoriatic, Antithyroid, Antiammatic, Antipsoriatic, Antithyroid, Antiammatic, Antiparkinsonian, Cerebroprotective, Vasotropic, Nootropic, Antiparkinsonian, Antiangiogenic, Antiarteriosclerotic, Virucide, Antibacterial, light chain variable region.
                                                                                Gaps
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                                                                                                                                                            61 RPSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                          Human IP10 antibody light chain variable region, SEQ ID NO 90.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Srinivasan M, Cardarelli JM, Wang
Lane TE, Keirstead HS, Liu MT;
                                                   Length 108;
                                                    Score 541; DB 4; Length 10
Pred. No. 3e-36;
3; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 35; SEQ ID NO 90; 179pp; English.
                                                                                                                                                                                                                                                       AEB01062 standard; protein; 108 AA
                                                    95.2%;
95.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-DEC-2004; 2004WO-US041506.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-DEC-2003; 2003US-0529180P
                                                                                                                                                                                                                                                                                                            (first entry)
                                                                              103; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Huang H,
Rangan VS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2005-467095/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    autoimmune diseases.
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                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB; AEB01088
                             Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2005058815-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Deshpande S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                            08-SEP-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Passmore D,
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  invention
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                                                                                                                                                                                                                                                                                  AEB01062;
                                                      Query Match
                                                                  Best Local
                                                                              Matches
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disease, Hashimoto's thyroiditis), Sjogren's syndrome, pulmonary inflammation (e.g., asthma, chronic obstructive pulmonary disease, pulmonary sarcoidosis, lymphocytic alveolitis), transplant rejection, spinal cord injury, brain injury (e.g., stroke), neurodegenerative diseases (e.g., Alzheimer's disease, Parkinson's disease), gingivitis, gene therapy-induced inflammation, diseases of angiogenesis, inflammatory kidney disease (e.g., IgA nephropathy, membranoproliferative atherosphritis, rapidly progressive glomerulonephritis), or atherosclerosis. It is also useful for treating a viral or bacterial infection involving unwanted IP-10 activity in a subject, where the viral infection is mediated by HIV, HCV, HSV-1 or SARS virus. The present sequence represents the amino acid sequence of a human IP10 monoclonal antibody light chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes an isolated human antibody (I) that specifically binds osteoprotegerin ligand (OPGI). Also described: (1) a pharmaceutical composition comprising a pharmaceutical carrier and a therapeutic amount of (I); (2) methods of treating an osteopenic disorder in a patient, comprising administering to a patient the pharmaceutical composition of (I) or a pharmaceutical amount of (I); and (3) a method for detecting OPGI in a biological sample, comprising contacting the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EIVIJTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated antibody that specifically binds osteoprotegerin ligand, useful for diagnosing or treating bone disorders, such as osteoporosis, bone loss from arthritis, Paget's disease or osteopenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human; antibody; osteoprotegerin ligand; OPGL; osteopenic disorder; osteopathic; antiarthritic; cytostatic; gene therapy; bone disorder; osteoporosis; bone loss; arthritis; Paget's disease; osteopenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16E1 anti-OPGL antibody kappa chain variable region SEQ ID NO:24.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPYTFGQGTKLEIK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 541; DB 9; Length 108;
Pred. No. 3e-36;
3; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95.2%;
95.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 95.4 Matches 103; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-PSDB; ADF11410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 108 AA;
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sample with (I) under conditions that allow for binding of the antibody to OPGL, and measuring the level of bound antibody in the sample. (I) has osteopathic, antiarthritic and cytostatic activities, and can be used in gene therapy. The composition and methods are useful in diagnosing or treating bone disorders, such as osteoporosis, bone loss from arthritis, baget's disease or osteopenia. The antibody (I) may also be used for detecting OPGL in biological samples and in identifying cells or tissues that produce the protein. The present sequence represents a sequence which is used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                      EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQXPGQAPRLLIYDASNRATGIPA
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                                                                                                                                                                                                                                                                                                                                                                                                                              Length 109;
                                                                                                                                                                                                                                           Score 541; DB 7; Length 10 Pred. No. 3.1e-36; 3; Mismatches 2; Indels
                                                                                                                                                                                                                                         95.2%;
95.4%;
                                                                                                                                                                                                                                                                                  Matches 103; Conservative
                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                         Sequence 109 AA;
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16El anti-OPGL antibody light chain SEQ ID NO:48
ADF11435 standard; protein; 215 AA
         (first entry)
         12-FEB-2004
    ADF11435;
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human; antibody; osteoprotegerin ligand; OPGL; osteopenic disorder; osteopathic; antiarthritic; cytostatic; gene therapy; bone disorder; osteoporosis; bone loss; arthritis; Paget's disease; osteopenia.

Homo sapiens

WO2003086289-A2.

23-OCT-2003

07-APR-2003; 2003WO-US010749.

05-APR-2002; 2002US-0370407P.

(AMGE-) AMGEN INC

Medlock B,

Boyle WJ,

New isolated antibody that specifically binds osteoprotegerin ligand, useful for diagnosing or treating bone disorders, such as osteoporosis, bone loss from arthritis, Paget's disease or osteopenia. WPI; 2003-845253/78. N-PSDB; ADF11434.

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Sullivan JK, Elliott RL, Martin F, Huang

English. Claim 16; SEQ ID NO 48; 156pp; The present invention describes an isolated human antibody (1) that specifically binds osteoprotegerin ligand (OPG1). Also described: (1) a pharmaceutical composition comprising a pharmaceutical carrier and a therapeutic amount of (1); (2) methods of treating an osteopenic disorder in a patient, comprising administering to a patient the pharmaceutical composition of (1) and (3) an enthod for detecting OPG1 in a biological sample, comprising contacting the sample with (1) under conditions that allow for binding of the antibody to OPG1, and measuring the level of bound antibody in the sample. (1) has osteopathic, antiarthritic and cytostatic activities, and can be used in gene therapy. The composition and methods are useful in diagnosing or

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, monoclonal antibody; antibody; interleukin-1; receptor;
antiasthmatic; antiinflammatory; dermatological; antiallergic;
protozoacide; antiinfleumatic; antiarthritic; osteopathic; vasotropic;
analgesic; antidiabetic; nephrotropic; antianaemic; nootropic;
anticonvulsant; dermatological; antigout; antiparkinsonian; antidiabetic;
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bone loss from arthritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anti-interleukin-1 receptor type 1 antibody kappa chain variable region.
                                                                                                                                                                                                                                 1 BIVLTOSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                            Gaps
                Paget's disease or osteopenia. The antibody (i) may also be used for detecting OPGL in biological samples and in identifying cells or tiss that produce the protein. The present sequence represents a sequence which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated human antibody that specifically binds interleukin-1 rectype 1 (IL-1R1) useful for treating IL-1 mediated diseases such srheumatoid arthritis, osteoarthritis and inflammatory conditions.
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                                                                                                                                                                                                                                                                                                 61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
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                                                                                                                                        Score 541; DB 7; Length 215;
Pred. No. 5.8e-36;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Martin F,
treating bone disorders, such as osteoporosis,
                                                                                                                                                                           3; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 12; 179pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                      ADM41547 standard; protein; 128
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                                                                                                                                        95.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-JUN-2004 (first entry)
                                                                                                                                                                           Matches 103; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2004-248462/23.
                                                                                                                                                            Similarity
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                                                                                                          Sequence 215 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytostatic.
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                                                                                                                                          Query Match
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eating-disorders-gen; immunomodulator; antimicrobial; cardiovascular-gen;
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                                                                                                                                                                                                                                                                                                                                                                   EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated mammalian anti-tumor necrosis factor (TNF) antibody, useful for diagnosing or treating an anti-TNF related condition, e.g. cancer, anorexia, cachexia, or bacterial infection.
                         sclenosis, Alzheimer's disease, cacheria, asthma, atthma, atthma, atthma, atthma, atthma, atthma, atthma, atthma, atthma associated illnesses, coronary conditions, cancer including leukaemia and tumour metastasis, diabetes, endometriosis, fever, fibromyalgia, glomerulonephritis, graft versus host disease, carcertritis, intemmatoid arthritis, inflammatory eye disease, ischaemia, Kawasaki's disease, learning impairment, lung disease, multiple sclerosis, myopathy, osteoporosis, pain, Parkinson's disease, periodontal disease, pre-term labour, psoriasis, reperfusion injury, septic shock, side effects of radiation therapy, temporal mandibular joint disease, sleep disturbance, uveitis, or an inflammatory condition resulting from strain, sprain, cartilage damage, trauma, orthopaedic surgery, infection or other disease processes.
               amyotrophic latera
                                                                                                                                                                                                                                                                                                                                       EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                                                                                                                          Gaps
 of IL-1R1 in a
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                                                                                                                                                                                                                                                                                                                                                                                                                  Length 128;
                                                                                                                                                                                                                                                                                                     3; Indels
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mediated diseases or for detecting the amount mediated diseases include acute pancreatitis,
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Pred. No. 4.3e-36;
1; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADD89877 standard; protein; 130
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/note= "CDR3"
                                                                                                                                                                                                                                                                         95.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 96.3
Matches 104; Conservative
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/note= "C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-804040/75.
N-PSDB; ADD89883.
                                                                                                                                                                                                                                              Sequence 128 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               neuroprotective.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003083061-A2
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The present sequence is that of the light chain variable region of human anti-tumour necrosis factor (TNF) monoclonal antibody 9F7. This human TNF reactive IgG monoclonal antibody was generated by cloning variable and constant region DNA in vector DC4 and expression in CHO cells. The invention provides isolated human, primate, rodent, mammalian, chimeric, humanized and/or CDR-grafted anti-TNF antibodies, immunoglobulins, their cleavage products, other specified portions and variants, as well as anti-TNF antibody compositions, nucleic acids encoding these, vectors, host cells, methods for producing the antibodies using a host cell, transgenic animal or transgenic plant or plant cell, and therapeutic compositions, methods and devices. The antibody, nucleic acid, protein, composition and methods are useful for diagnosing or treating an anti-TNF related condition, e.g. cancer, anorexia, cachexia, or an immune, cardiovascular, infectious, and/or neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tumour necrosis factor; TNF; immunotherapy; TNF related diseases; obesity; immune related disease; rheumatoid arthritis; cardiovascular disease; stroke; malignant disease; leuksemia; neurological disease; multiple solerosis; infection; hepatitis; anorectic; antiathritic; cerebroprotective; vasotropic; cytostatic; neuroprotective; antibacterial; antiinflammatory; hepatotropic; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95.1%; Score 540; DB 7; Length 130; 96.3%; Pred. No. 4.3e-36; ive 1; Mismatches 3; Indel8
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                                  Claim 1; Fig 3B; 87pp; English.
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(SCAL/) SCALLON B J.
(CART/) CARTON J M.
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The present invention relates to a mammalian anti-tumour necrosis factor (TNF) antibody capable of inhibiting binding of TNF alpha to TNF receptor. The invention is useful for diagnosing or treating an anti-TNF related condition in a cell, tissue, organ or animal and in immunotherapy. The invention is also useful for treating TNF related diseases chosen from obsesty, immune related diseases und as rheumatoid arthritis, cardiovascular disease such as stroke, malignant disease such as leukaemia, neurological disease such as stroke, malignant disease such as bacterial or viral infections such as hepatitis. The present sequence is the human 9FTLC light chain protein (variable region and J-region). This sequence is used in the exemplification of the invention.
                          Example 4; Fig 3; 45pp; English.
```

Sequence 130 AA;

Gaps .; 0 Length 130; 3; Indels 95.1%; Score 540; DB 8; 96.3%; Pred. No. 4.3e-36; iive 1; Mismatches 3; Query Match
Best Local Similarity 96.3
Matches 104; Conservative

셤

61 RFSGSGSGTDFTLIISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108 È

Search completed: December 28, 2005, 13:58:08 Job time : 128.462 secs

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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- protein search, using sw model OM protein

December 28, 2005, 13:46:34; Search time 22.1538 Seconds (without alignments) 469.057 Million cell updates/sec Run on:

US-09-920-137F-8 568 1 BIVLTQSPATLSLSPGERAT......QQRSNWPPFTFGPGTKVDIK 108

Perfect score: Sequence: **BLOSUM62** Scoring table:

Gapop 10.0 , Gapext 0.5

283416 segs, 96216763 residues Searched:

Total number of hits satisfying chosen parameters:

283416

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR 80:* Database

pir1:* pir2:* pir3:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	a kanna chain V	_	g kappa chain pr	kappa chain V	g kappa chain V-	b	ь	g kappa	ы	g kappa	kappa chain	g kappa chain	kappa chain	ы	kappa chain pr	g kappa chain	kappa chain V		kappa chain	kappa chain V-	kappa chain		kappa		kappa chain			kappa cha	
SUMMARIES	a		523628	PL0106	A56701	840345	G44151	S29627	840379	S57444	S40344	K3HUVG	JE0244	PC4282	S46375	S11697	S40362	S34005	840325	S40363	S40343	S40360	A30608	B26555	840328	JE0242	H30601	B30601	PH0963	S37506
	80		N	N	~	~	~	N	7	~	~	н	~	~	~	N	~	~	~	~	N	N	7	~	~	~	7	N	~	7
	* Query Match Length	: :	111	144	128	128	108	129	128	107	125	115	215	106	114	115	117	107	129	129	128	130	109	116	131	215	109	109	109	92
	Query Match		92.3	91.5	91.1	90.4	90.2	6.68	87.6	87.1	86.0	85.7	84.7	83.4	83.2	82.9	82.8	82.3	82.3	81.7	81.4	81.3	81.0	80.8				90.6		
	Score	52	524.5		517.5					494.5	488.5	487	481	473.5	472.5	471	470.5	467.5	467.5	464	462.5	462	460	459	459	459	458	458	458	456
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kappa	light	kappa	kappa												
Ig															- bi
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N	~	N	N	7	~	~	~	-	-	~	-	~	~	~	~
108	109	109	108	92	109	109	128	129	109	110	129	109	129	134	108
80.3	80.3	80.3	80.2	80.1	80.1	80.1	79.9	79.9	79.8	79.8	79.8	79.6	79.6	79.6	78.9
9	456	456	455.5	455	455	455	454	454	453	453	453	452	452	452	448
42															

ALIGNMENTS

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RESULT 1
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Ig kappa chain V region - human (fragment)

Ig kappa chain V region - human (fragment)

C; Species: Homo sapiens (man)

C; Accession: S54905

R; Esposito, G.; Traboni, C.

B; Esposito, G.

B; Esposito, C.

B; Esposito, G.

B; Esposito, C.

B; Esposi

ö Gaps ; 0 Score 528; DB 2; Length 114; Pred. No. 5e-39; 4; Mismatches 4; Indel8 4; Mismatches Query Match
Best Local Similarity 92.6%;
Matches 100; Conservative

61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108 셤 ò

Ig kappa chain V region - human (fragment) C;Species: Homo sapiens (man) C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 21-Jan-2000

C;Accession: S25628
R;Olee, T.; Lu, B.W.; Huang, D.F.; Soto-Gil, R.W.; Deftos, M.; Kozin, F.; Carson, D.A.;
R;Olee, T.; Lu, B.W.; Huang, D.F.; Soto-Gil, R.W.; Deftos, M.; Kozin, F.; Carson, D.A.;
A;Title: Genetic analysis of self-associating immunoglobulin G rheumatoid factors from t
A;Reference number: S23623; MUID:92156804; PMID:1740665
A;Accession: S23628
A;Accession: S23628
A;Accession: S23628
A;Accession: S23628
A;Accession: S23628

A; Residues: 1-111 <OLE>

A;Cross-references: UNIPARC:UP10000115F96; EMBL:X59705; NID:g34022; PIDN:CAA42226.1; PID C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin C;Keywords: heterotetramer; immunoglobulin F;16-90/Domain: immunoglobulin homology <IMM>

Length 111; DB 2; 92.3%; Score 524.5; Query Match us-09-920-137f-8.rpr

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EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
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Pred. No. 1e-37;
3; Mismatches 5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ig kappa chain V region (JM-10) - human (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Status: preliminary; translation not shown A;Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match

Best Local Similarity 91.7%;
Matches 100; Conservative 3
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G44151
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C;Species: Homo sapiens (man)
C;Species: Joban-1992 #sequence_revision 30-Jun-1992 #text_change 21-Jan-2000
C;Accession: PLO106
R;Silberstein, L.E.; Litwin, S.; Carmack, C.E.
J, Exp. Med. 169, 1611-1643, 1989
A;Title: Relationship of variable region genes expressed by a human B cell lymphoma secr
A;Reference number: PLO106; MUID:89235583; PMID:2541221
A;Accession: PLO106
A;Molecule type: mRNA
A;Residues: 1-144 «SIL»
A;Cross-references: UNIPARC:UP100001767A2
C;Superfamily: immunoglobulin V region; immunoglobulin
F;1-20/Domain: signal sequence #status predicted «SIG»
F;36-110/Domain: ummunoglobulin homology «IMM»
F;44-54/Region: complementarity-determining 1
F;10-115/Region: complementarity-determining 2
F;10-115/Region: complementarity-determining 3
F;116-127/Domain: J region (fragment) «CRE»
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19 kappa chain V region precursor (HuA) - human (fragment)

19 kappa chain V region precursor (HuA) - human (fragment)

19 kappa chain V region precursor (HuA) - human (fragment)

2,5pecies Home sapiens (man)

2,5pecies Home asform

3,500 Home 270, 1245-12465, 1995

4,7itle: Human and mouse monoclonal antibodies to blood group A substance, which are neally a preliminary

4,5pecies Home 270, 1245-12465, 1995

4,7itle: Human and mouse monoclonal antibodies to blood group A substance, which are neally sheetenen number: A56701

5,5tatus: preliminary

6,5tatus: preliminary

7,8tatus: mRNA

7,8ta
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                                                                                                                                     1 EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA 60
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                                                                                                                                                                                                                                                          Length 144;
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nes 3;
95.4%; Pred. No. 9.8e-39;
                              2; Mismatches
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ilarity 94.4%; Pred. No. 3.4e
Conservative 2; Mismatches
                                 Matches 103; Conservative
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   Best Local Similarity
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Best Local S:
Matches 102
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A;Cross-references: UNIPPARC:UP1000011615F; EMBL:X72455; NID:g441378; PIDN:CAA51123.1; PI CS.Superfeanily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin Pi32-106/Domain: immunoglobulin homology <IMM>
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R;Zebedee, S.L.; Barbas III, C.F.; Hom, Y.L.; Caothien, R.H.; Graff, R.; DeGraw, J.; Pya R;Zebedee, S.L.; Barbas III, C.F.; Hom, Y.L.; Caothien, R.H.; Graff, R.; DeGraw, J.; Pya Proc. Natl. Acad. Sci. U.S.A. 89, 3175-3179, 1992
A;Title: Human combinatorial antibody libraries to hepatitis B surface antigen. A;Reference number: A44151; MUID:92228746; PMID:1373487
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A;Note: nucleotide translation not given
C;Superfamily: immunoglobulin vegion; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-89/Domain: immunoglobulin homology <IMM>
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Cispecies: Homo sapiens (man)
Cispecies: Homo sapiens (man)
Cipate: 19-May-1994 #sequence_revision 26-May-1995 #text_change 21-Jan-2000
CiAccession: $40345
Bur. J. Immunol. 23, 3248-3271, 1993
A;Title: Expressed human immunoglobulin chi genes and their hypermutation.
A;Reference number: $40312; MUID:94080891; PMID:8258341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Species: Homo sapiens (man)
C;Date: 27-Jun-1994 #sequence_revision 27-Jun-1994 #text_change 21-Jan-2000
                                       21 BIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA
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                                                                                                                                                                            RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPR-SFGQGTKVEIK 127
                                                                                                                                        61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
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A; Status: preliminary; translation not shown
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Best Local Similarity 90.64
Matches 96; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 97; Conserv
                                                                                                                                                                                                        A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-107 <PAT>
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A; Residues: 1-125 < KLE>
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RESULT 7
S29627
IQ Wappa chain V region (60.3 hybridoma) - human
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 21-Jan-2000
C;Accession: 334110; $29627
R;Walls, M.A.; Hsiaco, K.; Harris, L.J.
Nucleic Acids Res. 21, 2921-2929, 1993
A;Title: Vectors for the expression of PCR-amplified immunoglobulin variable domains wit A;Reference number: S34110; MUD:99324379; PMID:8932501
A;Accession: S3410
A;Accession: S3410
A;Accession: Gid sequence not shown; translation not shown
A;Molecule type: mRNA
A;Residues: 1-129 «MA2>
A;Cross-references: UNIPARC:UP1000011649A; EMBL:Z17330; NID:938511; PIDN:CAA78978.1; PIC
A;Note: human sequences cloned and sequenced prior to expression in mouse myeloma cells
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, October 1992
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P
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Grappa chain Ray-1994 #sequence_revision 26-May-1995 #text_change 21-Jan-2000

Gracession: $40379

Br.Klein, R.; Jaenichen, R.; Zachau, H.G.

Eur. J. Immunol. 23, 3248-3271, 1993

Ariticle: Expressed human immunoglobulin chi genes and their hypermutation.

Areference number: $40312; MUID:94080891; PMID:8258341

Areference number: $40379

Arecession: $40379

Areference number; translation not shown

Areference type: mRNA

Areference: UNIPARC:UPI0000116181; EMBL:X72489; NID:9441446; PIDN:CAA51157.1; If Cranslatinity: immunoglobulin V region; immunoglobulin homology

Grappa creference: immunoglobulin homology <IMM>

From the credit immunoglobulin homology <IMM>
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C;Keywords: heterotetramer; immunoglobulin
F;36-110/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     atch 89.9%; Score 510.5; DB 2; cal Similarity 92.7%; Pred. No. 1.8e-37; 101; Conservative 1; Mismatches 6;
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Ig kappa chain V-J region - human (fragment)
C;Species: Homo sapiens (man)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Introns: 17/1
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Matches 10
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C; Date: 10-Oct-1995 #sequence_revision 17-Nov-1995 #text_change 21-Jan-2000 C; Accession: S57444 B. R; Paterson, G.; Wilson, G.; Kennedy, P.G.E.; Willison, H.J. Bubmitted to the EMBL Data Library, June 1995 A; Description: Analysis of anti-GMI ganglioside IGM antibodies cloned from motor neuropa A; Reference number: S57408 A; A; A; Cession: S57444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross_references: UNIPARC:UP1000011615B; EMBL:X72454; NID:g441376; PIDN:CAA51122.1; PIC;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Accession: A01900
R;Pech, M.; Zachau, H.G.
Nucleic Acids Res. 12, 9229-9236, 1984
A;Title: Immunoglobulin genes of different subgroups are interdigitated within the V-K 1 A;Reference number: A93549; MUID:85087932; PMID:6440122
A;Accession: A01900
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A;Residues: 1-115 <PEC>
A;Cross-references: UNIPROT:P04433; UNIPARC:UPI0000116D05; GB:X01668; GB:K02768; NID:933
                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: UNIPARC:UDI0001137AE; EMBL:X87898; NID:g871275; PIDN:CAA61149.1; C;Superfamily: immunoglobulin V region; immunoglobulin homology F;16-90/Domain: immunoglobulin homology <IMM>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ig Kappa chain V-J region - human
C;Species: Homo sapiens (man)
C;Date: 19-May-1994 #sequence_revision 26-May-1995 #text_change 21-Jan-2000
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C;Species: Homo sapiens (man)
C;Date: 17-Mar-1987 #sequence_revision 17-Mar-1987 #text_change 09-Jul-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Accession: §40344
R;Klein, R.; Jaenichen, R.; Zachau, H.G.
R:L. J. Immunol. 23, 3248-3271, 1993
A;Title: Expressed human immunoglobulin chi genes and their hypermutation. A;Reference number: $40312; MUID:94080891; PMID:8258341
A;Accession: $40344
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 107;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              87.1%; Score 494.5; DB 2;
89.8%; Pred. No. 3.7e-36;
7ative 3; Mismatches 7;
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Pred. No. 1.4e-35;
3; Mismatches 6
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A; Reference number: PC4279; MUID: 97236289; PMID: 9125110
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                                                                                                                                                             A; Accession: PC4284
A; Molecule type: protein
A; Residues: 1-106 <SU2>
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A; Residues: 1-114 <BEN>
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A;Note: the sequence was determined from the germline gene C;Genetics:
A;Gene: GDB:IGKV3
A;Cross-references: GDB:136266
A;Map position: 2p12-2p1
A;Introns: 17/1
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kap hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into la C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin v-III region (Vg) #status predicted <NAT>
F;12-00Domain: signal sequence #status predicted <SIG>
F;21-115/Product: Ig kappa chain V-III region (Vg) #status predicted <MAT>
F;21-13/Region: framework 1
F;55-69/Region: complementarity-determining 1
F;55-69/Region: complementarity-determining 2
F;70-76/Region: complementarity-determining 3
F;109-115/Region: complementarity-determining 3
F;43-108/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JE0244

1g kappa chain NIG2 precursor - human

G;Species: Homo sapiens (man)

C;Species: Homo sapiens (man)

C;Date: 05-Dec-1998 #text_change 21-Jan-2000

C;Accession: JE0244

R;Alim, M.A.; Hara; Y.; Hossain, M.S.; Takeda, K.; Yamagata, F.; Yamaki, S.; Kazi, H.; T

submitted to JIPID, November 1998

A;Description: A new subgroup of k type light chains (VkV) identified in cases of AL amy

A;Reference number: JE0244

A;Residues: 1-215 <ALI>
A;Residues: 1-215 <ALI>
C;Superfamily: immunoglobulin V region; immunoglobulin homology

F;16-90/Domain: immunoglobulin homology <INM>
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PC4282
Ig Asapa chain (anti-SS-A/Ro 60K peptide) (E-42 and E-56) - human (fragment)
Ig Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 07-Jul-1997 #sequence_revision 29-Aug-1997 #text_change 21-Jan-2000
C;Date: 07-Jul-1997 #sequence_revision 29-Aug-1997 #text_change 21-Jan-2000
C;Date: 07-Jul-1997 #sequence_revision 29-Aug-1997
R;Suzuki, H.; Takemura, H.; Suzuki, M.; Sekine, Y.; Kashiwagi, H.
Biochem. Biophys. Res. Commun. 232, 101-106, 1997
A;Title: Molecular cloning of anti-ss-A/Ro 60-kDa peptide fab fragments from infiltratin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 115;
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Pred. No. 1.8e-35;
0; Mismatches 1; Indels
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85.2%; Pred. No. 1.1e-34;
iive 7; Mismatches 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWP 95
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Best Local Similarity 98.9%;
Matches 94; Conservative
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Matches 9
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A,Note: E-56
C,Comment: This antibody is commonly found in systemic autoimmune diseases such as Sjoeg
C,Superfamily: immunoglobulin V region; immunoglobulin homology
F;14-88/Domain: immunoglobulin homology <IMM>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 25-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
C; Accession: 811697 # Sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
C; Accession: 811697 # Sequence_revision 10-Nov-1995 # Secondary # C; Accession: 811697 # WUID: 85264787; PMID: 3927006
A; Accession: 811697
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EMBO J. 13, 2951-2962, 1994
A;Title: Human lupus anti-DNA autoantibodies undergo essentially primary V(chi) gene A;Reference number: $46369; MUID:94313975; PMID:8039491
A;Accession: $46375
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C'Species: Homo sapiens (man)
C'Date: 27-Jan-1995 #sequence_revision 01-8ep-1995 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 VLTQSPATLSLSPGERATLSCRASQSVSNYLAWYQQKPGQAPRLLIYDISKRATGIPAKF
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85.3%; Pred. No. 3.2e-34;
live 6; Mismatches 9;
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85.8%; Pred. No. 2.4e-34;
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A, Molecule type: protein
A, Residues: 1-106 <SUZ>
A, Cross-references: UNIPARC:UP10000176E92
A, Note: E-42
                                                                                                                                                                                                                                                   A; Cross-references: UNIPARC: UPI0000176E92
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A;Cross-references: UNIPARC:UPI0000115E41; EMBL:X17264; NID:g37898; PIDN:CAA35168.1; PID A;Note: the nucleotide sequence was submitted to the EMBL Data Library, December 1989 C;Genetics:
A;Introns. 17/1
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;36-110/Domain: immunoglobulin homology <IMM>
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0; Gaps Query Match
82.9%; Score 471; DB 2; Length 115;
Best Local Similarity 96.8%; Pred. No. 4.3e-34;
Matches 91; Conservative 0; Mismatches 3; Indels

ö 1 EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA 60

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Search completed: December 28, 2005, 14:04:03 Job time : 23.1538 secs

िंड Paga Blank (uspto)

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85.7%; Score 487; DB 1; Length 115; llarity 98.9%; Pred. No. 9.5e-43; Conservative 0; Mismatches 1; Indels
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                                       ; Search time 137.077 Seconds (without alignments) 555.871 Million cell updates/sec
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Copyright (c) 1993 - 2005 Compugen Ltd.
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                                                                                                                2166443 segs, 705528306 residues
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This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
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Pech M., Zachau H.G.;
"Immunoglobulin genes of different subgroups are interdigitated within
the VK locus.";
                                                                                                                                                                                                                                                                                                                                                       13-AUG-1987 (Rel. 05, Created)
13-AUG-1987 (Rel. 05, Last sequence update)
10-MAY-2005 (Rel. 47, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
1g kappa chain V-III region VG precursor (Pragment).
Homo sapiens (Human).
Eukaryota, Metacon, Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R EMBL; X01668; -; NOT_ANNOTATED_CDS; Genomic_DNA.
R MSP; P01905; M3HUVG.
R HSSP; P01625; 1BEQ.
R GO; GO:0005925; F: antigen binding; NAS.
R GO; GO:0006925; P: fantigen binding; NAS.
R GO; GO:0006925; P: fantigen binding; NAS.
R InterPro; IPR007110; Ig-like.
R InterPro; IPR003596; Ig-v.
R SWART; SW00406; IG-v.
R SWART; SW00406; IG-v.
R SWART; PS50835; IG_LIKE; 1.
R INDENDOJOULIN GOMBIN; IMMUNOGLOBULIN V region; Signal.
Q9ul77 P
P01600 P
Q56919 P
Q9ul81 P
P01663 P
P01603 P
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              KV1H HÜMAN
OS6915 HÜMAN
CO9UL81 HÜMAN
KV3M MOUSE
KV4C HÜMAN
KV1K HÜMAN
KV1L HÜMAN
KV3D MOUSE
OSGWÜL HÜMAN
KV3L HÜMAN
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KV3H MOUSE
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Complementarity-determining-2. Framework-3. Complementarity-determining-3.

12575 MW; 2DE47CDA3A17D555 CRC64;

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PROSITE; PS50835; IG LIKE; 1.
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Wagner S.D., S.J., S.J.,
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                           EIVLTQSPATLSLSESPGERATLSCRASQSVSSYLAWYQQKPQQAPRLLIYDASNRATGIPA
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                     EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Myosin-reactive autoantibodies in rheumatic carditis and normal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531; Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
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01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Myosin-reactive immunoglobulin light chain variable region
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88.0%; Pred. No. 5.3e-42;
ive 5; Mismatches 7;
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                                                                                                                                                                                                                                                                                         01-MAY-2000 (TrEMBLrel. 13, 01-MAY-2000 (TrEMBLrel. 13,
                                                                                                                                                                                                                                                    Q9UL83_HUMAN PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PubMed=8436174;
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ID KV3H HUMAN
AC P04207;
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Best Local
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20-MAR-1987 (Rel. 04, Created)
01-NOV-1990 (Rel. 16, Last sequence update)
10-MAY-2005 (Rel. 47, Last amontation update)
10-MAY-2005 (Rel. 47, Last amontation update)
11 kappa chain V-III region CLL precursor (Rheumatoid factor).
12 kappa chain W-III region CLL precursor (Rheumatoid factor).
13 kappa chain W-III region CLL precursor (Rheumatoid factor).
14 kmom Sapiens (Human).
15 kmom M-III kmom W-III k
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Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi;
Mammalia, Butheria, Euarchontoglires, Primates, Catarrhini, Hominidae;
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MEDLINE=66177570; PubMed=3083417;
Jirik P.R., Sorge J., Fong S., Heitzmann J.G., Curd J.G., Chen P.P.,
Goldfien R., Carson D.A.,
Cloning and sequence determination of a human rheumatoid factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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Complementarity-determining-3.
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1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-CCT-2003 (TrEMBLrel. 25, Last annotation update)
Myosin-reactive immunoglobulin kappa chain variable region
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Pred. No. 2.5e-41;
9; Mismatches 9; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunoglobulin domain; Immunoglobulin V region; Signal.
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Proc. Natl. Acad. Sci. U.S.A. 83:2195-2199(1986).
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01-NOV-1990 (Rel. 16, Last sequence update)
01-NOY-1990 (Rel. 47, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
1g kappa chain V-III region HAH precursor.
Homo sapiene (Human)
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 BIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
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MEDLINE=88171307; PubMed=3127527; DOI=10.1084/jem.167.3.840;
Kipps T.J., Tomhave E., Chen P.P., Carson D.A.;
Autoantibody-associated kappa light chain variable region gene expressed in chronic lymphocytic leukemia with little or no somatic mutation. Implications for etiology and immunotherapy.";
J. Exp. Med. 167:840-852(1988).
-1- DISEASE: The protein is one of the surface immunoglobulin M autoantibodies expressed in patients with chronic lymphocytic
                                                                                                                                             "Myosin-reactive autoantibodies in rheumatic carditis and normal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .
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                                                                             MEDLINE=99277139; PubMed=9614934; DOI=10.1006/clin.1998.4531; Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M., Young D.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 109;
                                                                                                                                                                                                                                                                                                                                                                                                                                        82.4%; Score 468; DB 2; Lengua 2.82.4%; Pred. No. 8.5e-41; Lindels
                                                                                                                                                                                                                                                                                                                                                                                                                 109 AA; 11761 MW; FB1E43E7C7AFACCC CRC64;
                                                                                                                                                                               Cin. Immunol. Immunopathol. 87:184-192(1998).

EMBL; AF035029; AAD56265.1; -; mRNA.

PIR; D30609; D30609.

HSSP; P01625; 1EK3.

SMR; Q9UL85; 1-109.

InterPro; IPR003110; Ig-like.

InterPro; IPR003596; Ig_v.

SMART; SM00406; IGv; 1.
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HSSP; P01625; 1EEQ.
SMR; P18135; 21-129.
Ensembl; ENSG00000165769; Homo sapiens.
GO; GO:0005576; C:extracellular region; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
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                                                              NUCLEOTIDE SEQUENCE
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Homo.
NCBI_TaxID=9606;
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NCBI_TaxID=9606;
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P18135;
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KV3L HUMAN
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use as long as its content is in no way modified and this statement is not
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Mammalia; Butheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=76276460; PubMed=60899; Klapper D.G., Capra J.D.; Klapper D.G., Capra J.D.; "The amino acid sequence of the variable regions of the light chains from two idiotypically across reactive IgM anti-gamma globulins."; Ann. Immunol. (Paris) 1270:261-271(1976).
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                                                                                                                                             chain V-III region HAH.
                                                                                                                                                                                              Complementarity-determining-1.
                                                                                                                                                                                                                                              Complementarity-determining-2.
                                                                                                                                                                                                                                                                     Framework-3.
Complementarity-determining-3.
                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 454; DB 1; Length 129,
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                                                                                        Immunoglobulin domain; Immunoglobulin V region; Signal. SIGNAL 1 20 Ig kappa chain V-III regio
                                                                                                                                                                                                                                                                                                                                                                                         14073 MW; D3C55292772774D0 CRC64;
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PROSITE; PS50835; IG LIKE; 1.
Direct protein sequencing; Immunoglobulin domain;
DISULFID 23 89
By similarity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-NAY-2005 (Rel. 47, Last annotation update)
Ilg kappa chain V-III region POM.
Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. 3e-39;
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SMR; PO1624; 1-109.
GO; GO:0005576; C:extracellular region; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-11ke.
InterPro; IPR003596; Ig-V.
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                                                                                                                                             Ig kappa cha
Framework-1
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InterPro; IPR007110; Ig-11ke.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
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                                                                                                                                                                                                                                                                                                                                                                                            129 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antigen.
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               Matches
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PIR;
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                                                                                                                                1 BIVMTQSPVTLSVSPGERATLSCRASQSISNSYLAWYQQKPSGSPRLLIYGASTRATGIP 60
                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                    1 EIVLTQSPATLSLSPGERATLSCRASQSV-YSYLAWYQQKPGQAPRLLIYDASNRATGIP
                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLING=88171307; FubMed=3127527; DOI=10.1084/jem.167.3.840; Kipps T.J., Tomhave E., Chen P.P., Carrson D.A.; "Autoantibody-associated kappa light chain variable region gene expressed in chronic lymphocytic leukemia with little or no somatic mutation. Implications for etiology and immunotherapy."; J. Exp. Med. 167:840-852(1988).

-1- DISEASE: The protein is one of the surface immunoglobulin M autoantibodies expressed in patients with chronic lymphocytic
                                                                         2;
                                                                                                                                                                            60 ARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
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Pramework-1.
                                            Length 109;
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                                                                       9; Indels
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SMART, SMORGE, IGY; 1.
PROSITE; PSS0835; IG_LIKE; 1.
Immunoglobulin domain; Immunoglobulin V region; Signal.
               62821DDC6A8ABA86 CRC64;
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                                          Score 453; DB 1;
Pred. No. 3.1e-39;
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HSSP; P01625; 1BEQ.
SMR; P18136; 21-129.
Ensembl; ENSG0000169769; Homo sapiens.
GO; GO:00038576; C:extracellular region; NAS.
GO; GO:0003857; P:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR07110; Ig-like.
InterPro; IPR007110; Ig-like.
                                                                                                                                                                                                                                                                                           01-NOV-1990 (Rel. 16, Created)
01-NOV-1990 (Rel. 16, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
How bappa chain V-III region HIC precursor.
                                                                                                                                                                                                                                                                 129 AA
                                                                       9; Mismatches
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               109 AA; 11922 MW;
                                         79.8%;
                                                        Local Similarity 81.7
hes 89; Conservative
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43
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ID KV3M HUMAN
AC P18136;
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Length 129;

79.8%; Score 453; DB 1;

Query Match

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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                        21 EIVLTQSPGTLSLSPGERATLSCRASQSVSSSYLAWYQQKPGQAPRLLIYGASSRATGIP
                                        BIVLTQSPATLSLSPGERATLSCRASQSV-YSYLAWYQQKPGQAPRLLIYDASNRATGIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Magner S.D., Luzzatto L.,
"V kappa gene segments rearranged in chronic lymphocytic leukemia are
distributed over a large portion of the V kappa locus and do not show
               Gaps
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               2,
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                                                                                                              60 ARPSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-98277139; PubMed-9614934; DOI=10.1006/clin.1998.4531; Mu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PubMed=1552291;
Martin T., Duffy S.F., Carson D.A., Kipps T.J.;
"Evidence for somatic selection of natural autoantibodies.";
J. Exp. Med. 175:983.991(1992).
                                                                                                                                                                                                                        01-MAY-2000 (TrEMBLrel. 13, Created)
U-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin light chain variable region
              Indels
              10;
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85.3%; Pred. No. 3.8e-39;
tive 4; Mismatches 10
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                                                                                                                                                                                           Q9UL78 HUMAN PRELIMINARY;
Q9UL78;
              93; Conservative
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01-MAY-2000
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H30607;
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KV3D HUMAN
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Biochemistry 20:5816-5822(1981).
-!- MISCELLANBOUS: This chain was isolated from an IgM with anti-gamma
globulin activity.
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Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Andrews D.W., Capra J.D.; "Amino acid sequence of the variable regions of light chains from two idiotypically cross-reactive human IgM anti-gamma-globulins of the Wa
                                                                                                                                                                                                                                                                                                                                                                                                        2; Gaps
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                                                                                                                                                                                                                                                                                                                                                            Length 109;
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85.3%; Pred. No. 8.1e-39;
ive 3; Mismatches 11; Indels
                                                                                                                                                                                                                                                                                                                  109 AA; 11646 MW; 5F675C52EC7EE197 CRC64;
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PROSITE; PS50835; IG LIKE; 1.
Direct protein sequencing; Immunoglobulin domain;
Immunoglobulin V region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
IG Mappa chain V-III region SIE.
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SMR; P01620; 1-109.
GO; GO:0005876; C:extracellular region; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006855; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR00396; Ig_v.
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                                                                              PIR, PH0965; PH0965.
PIR, S33988; S33988.
PIR, S34096, S34096.
HSSP; PO1625; 1EK3.
SMR; Q9UL78; 1-109.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig-v.
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Best Local Similarity 85.39
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Suter L., Barnikol H.U., Watanabe S., Hilschmann N.;
"Rule of antibody structure. The primary structure of a monoclonal immunoglobulin L-chain of kappa-type, subgroup 3 (Bence-Jones protein Ti). IV. The complete amino acid sequence and its significance for the mechanism of antibody production.";
Hoppe-Seylar's Z. Physiol. Chem. 353:189-208(1972).
-!- MISCELLANBOUS: The C region of this chain has the INV (3) marker.
-!- MISCELLANBOUS: This is a Bence-Jones protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SMART; SM00406; IGV; 1.-
PROSITE; PS50835; IG LIKE; 1.
Bence-Jones protein; Direct protein sequencing; Immunoglobulin domain;
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Mammalia, Eutheria, Euarchontoglires, Primates, Catarrhini, Hominidae,
                                                                                                                                                                                                    EIVITOSPATISISPGERATISCRASOSV-YSYLAWYQQKPGQAPRLLIYDASNRATGIP
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                                                        78.9%; Score 448; DB 1; Length 109;
                                                                                                              10; Indels
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109 AA; 11775 MW; 7689C3ECD646FFB4 CRC64;
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21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Ig kappa chain V-III region Ti.
Homo sapiens (Human).
                                                                                   Pred. No. 1e-38;
5; Mismatches
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SMR; P01622, 1-109.
GO; GO:0005576; C:extracellular region; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-11ke.
InterPro; IPR03596; Ig-V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         109 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              By similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PROTEIN SEQUENCE.
MEDLINE=72188439; PubMed=5027703;
                                                                                      84.48;
                                                  Query Match
Best Local Similarity 84.4
Matches 92; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunoglobulin V region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIR; A01895; K3HUTI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          92;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KV3D HUMAN
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RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Raha S.S., Logdelland N.B., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S.W., McKernan K.J., Malek J.A., Gunarathe P.H.,
RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C.,
RA Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
R Schmerch A., Schein J.E., Jones S.J.M., Marra M.A.,
R Generation and initial analysis of more than 15,000 full-length human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hypothetical protein.
Hypothetical protein.
Homo sapiens (Human).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Mammalia, Eutheria, Euarchontoglires, Primates, Catarrhini, Hominidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 EIVLTQSPATLSLSPGERATLSCRASQSV-YSYLAWYQQKPGQAPRLLIYDASNRATGIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21 ENVIJOSPGTISLSPGERATISCRASOSISSSYLAWYOOKPGOAPRILIYGVSSRATGIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE-Lung;
MEDLINE-22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
MEDLINE-22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
Strausherg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
Altechul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
Hopkins R.F., Ozordan H., Moore T., Max S.I., Wang J., Heileh F.,
Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            81 DRFSGSGSGTDFTLTISRLEPEDFAVYYCQQYGTSRPITFGQGTRLDIK 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 60 ARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                    TECHNEL Brain; Strausberg R.; Strausberg R.; Submitted (JUN-2002) to the EMBL/GenBank/DDBJ databases. EMBL; BC03451; AAH32451.1; -; mRNA. HSSP; PO1837; IKCU.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PROSITE; PS50835; IG_LIKE; 2.
PROSITE; PS02290; IG_MHC; UNKNOWN_1.
Hypothetical protein.
SEQUENCE 236 AA; 25834 MW; 6647A9E77A3C0053 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                          Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          78.8%; Score 447.5; DB 2
82.6%; Pred. No. 2.9e-38;
iive 4; Mismatches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SMC, Q6PLGB; 21-236.
InterPro; IPR003599; IG.
InterPro; IPR003599; IG-like.
InterPro; IPR003509; IG-like.
InterPro; IPR003006; IG-MHC.
InterPro; IPR003596; IG-W.
Fam; Pr007654; C1-8et; I.
SMART; SM00409; IG; 2.
SMART; SM00407; IGC1; I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             QEPJF2 HUMAN PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 82.6 es 90; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUCLEOTIDE SEQUENCE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q6PJF2;
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                                                                                                                                                                                                                                                                                  (Fragment).
Homo sapiens (Human).
Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypothetical protein.
Hypothetical protein.
Homo sapiens (Human).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Euarchontoglires, Primates, Catarrhini, Hominidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 EIVLTQSPATLSLSPGERATLSCRASQSV-YSYLAWYQQKPGQAPRLLIYDASNRATGIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
Klausner R.D., Collins F.S., Wagner L., Shemmen C.M., Schuler G.D.,
Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Myosin-reactive autoantibodies in rheumatic carditis and normal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DRFSGSGSETDFTLTISRLEPEDFAVYYCQQYGS-SIFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            60 ARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                        61 DRFSGSGSGTDFTLTISRLEPEDFAVYYCQQYGS-SPSTFGQGTKVELK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-98277139; PubMed-9614934; DOI=10.1006/clin.1998.4531; Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  78.9%; Score 448; DB 2; Length 109; llarity 85.3%; Pred. No. 1e-38; Conservative 2; Mismatches 12; Indels
                                                                                                                                                                                                       01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Myosin-reactive immunoglobulin kappa chain variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       109 AA; 11928 MW; 243325F72C7DAC83 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Clin. Immunol. Immunopathol. 87:184-192(1998).
EMBL; AF035028; AAD56264.1; -; mRNA.
PIR; B30607; B30607.
PIR; I30601; 130601.
PIR; P01625; 1EK3.
SMR; Q9UL86; 1-109.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ensembl; ENSG0000169769; Homo sapiens.
InterPro; IPR007110; Ig-11ke.
InterPro; IPR003196; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
                                                                                                                                                                                         Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q6PIL8;
05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
                                                                                                                                                                                      01-MAY-2000 (TrEMBLrel. 13, 01-MAY-2000 (TrEMBLrel. 13,
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QEPIL8_HUMAN PRELIMINARY;
                                                                                                                                         Q9UL86_HUMAN PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=9606;
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NON TER
SEQUENCE
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                                                                                                                     HUMAN
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RESULT 1

OGUISS 6

HOTO COST 1

OGUISS 6

OGU

RESULT 12
OGPILIA HUM
OG PILIA HUM
OG PILIA
DT 05-JU
DT 05-JU
DT 05-JU
DE HYPOT
OS HOMO
OC MAMMA
OC HOMO.
OC HOMO.
OC HOMO.
RN (1)
RN (1)
RN (1)
RN TISSU
RX MEDLII
RA KLAUS
RA KLAUS
RA ALLSO
RA HOPKI

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59 80

Gaps

1;

14; Indels

Length 236;

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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Boak S.A., McEwan P.J., Malek J.A., Gunaratne P.H., Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., Villalon D.K., Muzny D.M., Sodergren E.J., Iu X., Gibbs R.A., Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A., Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C., Schmitz J., Myers R.M., Schnerch A., Schmitz J., Myers R.M., Schnerch A., Schein J.E., Schmetz J., Skalska U., Smallus D.E., Schnerch A., Schein J.E., Jones B.J.M., Marra M.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G6P5S8;
05-UUL-2004 (TrEMBLrel. 27, Created)
05-UUL-2004 (TrEMBLrel. 27, Last sequence update)
05-UUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein.
Hypothetical protein.
Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Ruarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DRFSGSGSGTDFTLTISRLEPEDFAVYYCQQYGS-SQGTFGPGTKVDIK 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          60 ARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Glandular pool- thyroid;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 235;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         78.7%; Score 447; DB 2; Length 23 84.4%; Pred. No. 3.3e-38; ive 5; Mismatches 10; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tisoublerg R.;
Straubberg R.;
Straubberg R.;
Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; BC016380; AAH16380.1; -; mRNA.
HSSP; P01837; 1KCU.
SNR; Q6PJF2; 21.235.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11 procein.
235 Aa; 25520 MW; F33A145A396BA285 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SWR, QGPDE2, 21-235.
INTERPRO, IRR003599, IG-like.
INTERPRO, IPR003599, IG-like.
INTERPRO, IPR003599; IG-like.
INTERPRO, IPR003006; IG-MHC.
INTERPRO, IRR003596; IG-MHC.
SWART; SW00409; IG; 2.
SWART; SW00409; IG; 2.
FROSITE; PSC0859; IG_LIKE; 2.
PROSITE; PSC0859; IG_LIKE; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                     and mouse cDNA sequences.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         QEP5S8_HUMAN PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          92; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUCLEOTIDE SEQUENCE.
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NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hypothetical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
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AC 066958
DT 05-JU
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Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H., Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A., Hulyk S.W., Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A., S.W., Witting M., Madan A., Rodrigues S., Sanchez A., Mhiting M., Touchman J.W., Green B.D., Dickson M.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Schein J.S., Now Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.B., Jones S.J.M., Marra M.A., "Generation and initial analysis of more than 15,000 full-length human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo Bapiens (Human).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Butheria, Buarchontoglires, Primates, Catarrhini, Hominidae,
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Andrews D.W., Capra J.D.;
"Amino acid sequence of the variable regions of light chains from two
idiocypically cross-reactive human IgM anti-gamma-globulins of the Wa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Strausberg R.; Submitted (NOV-2003) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hypothetical protein.
SEQUENCE 236 AA; 25773 MW; 953B37BEB4FF5F27 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                             Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
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Pred. No. 7.6e-38;
9; Mismatches 12
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21-JUL-1986 (Rel. 01, Last sequence update)
10-MXY-2005 (Rel. 47, Last annotation update)
1g kappa chain V-III region WOL.
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SWR; QCPSS8; 21-236.
InterPro; IPR007139; IG.
InterPro; IPR007109; IG-11ke.
InterPro; IPR007109; IG-11.
InterPro; IPR0073597; IG-1.
InterPro; IPR0073597; IG-1.
SWART; SW00409; IG; 2.
SWART; SW00409; IG; 2.
SWART; SW00409; IG; 2.
FROSTER; PS50835; IG LIKE; 2.
PROSTER; PS50835; IG LIKE; 2.
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Local Similarity 79.8%;
nes 87; Conservative 9
                                                                                                                                                                                                                                                                                                                                                                                               and mouse cDNA sequences.
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PROTEIN SEQUENCE.
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This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bioinformatics Institute. There are no restrictions on its use as long as its content is in to way modified and this statement is not
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 109;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 77.8%; Score 442; DB 1; Length 10 Best Local Similarity 84.4%; Pred. No. 4.3e-38; Matches 92; Conservative 3; Mismatches 12; Indels
                                                                                                                                                                                                                                                                                                                                                                 23 89 By similarity.
109 109
109 AA, 11746 MW, 566C115E6B9CBEBE CRC64;
                                                                                                              PIR; A01896; K3HUWL.
HSSP; P01625; 1LVE.
SNR; P01625; 1LVE.
SNR; P01625; 1LVE.
GO: GO: 0003823; 1-109.
GO: GO: 0003823; F: antigen binding; NAS.
GO: GO: 0005956; C: extracellular region; NAS.
GO: GO: 0005956; P: immune response; NAS.
INTERPRO; IPR003196; Ig-1ike.
INTERPRO; IPR003596; Ig-1ike.
SNART; SM00406; IGV; 1.
BY SNART; SM00406; IGV; 1.
Direct porctein sequencing; Immunoglobulin domain;
Immunoglobulin V region.
DISQUEND 23 89 By similarity.
NON TER 109 AA; 11746 MW; 566C115E6B9CBEEE CRC64
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Search completed: December 28, 2005, 14:03:09 Job time : 138.077 secs

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SQUENCE 46, Application US/09453234
Patent No. 6794132
GENERAL INFORMATION:
APPLICANT: Buechler, Joe
APPLICANT: Unbergy, Jeff
APPLICANT: Lonbergy, Nila
APPLICANT: GenPharm Incernational
TITLE OF INVENTION: Human Antibodies
FILE REFERENCE: 0220015-000110US
CURRENT APPLICATION NUMBER: US/09/453,234;
CURRENT FILING DATE: 1999-12-01
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86, Appli
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(without alignments)
297.632 Million cell updates/sec
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/cgn2_6/ptodata/1/iaa/H_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
/cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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US-09-453-234-80
US-09-453-234-90
US-09-456-090A-36
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US-09-456-090A-36
US-09-456-090A-36
US-09-456-090A-38
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US-09-456-090A-72
US-09-453-234-72
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                                                                                                                                                                                                                                                                                                                 572060 seqs, 82675679 residues
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Maximum Match 100%
Listing first 45 summaries
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Sequence 46, Application US/09456090A

Patent No. 6680209

GENERAL INFORMATION:

APPLICANT: Buechler, Joe

APPLICANT: Cray, Jeffs Genare

APPLICANT: Lonberg, Nils

TILLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS

TILLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS

TILLE OF INVENT APPLICATION UNDERS: US/09/456,090A

CURRENT FILING DATE: 1999-12-06

NUMBER OF SEQ ID NOS: 110

SOFTWARE: Patentin Ver. 2.1

LENGTH: 224
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US-09-490-070A-47
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US-09-490-153-47
US-09-456-090A-50
US-09-456-090A-86
US-09-456-090A-86
US-09-456-090A-86
US-09-453-234-86
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US-09-453-234-86
US-09-453-234-86
US-09-453-234-86
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US-09-453-234-86
US-09-453-234-86
US-09-473-087-74
US-09-472-087-17
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; ORGANISM: Homo sapiens
; OTHER INFORMATION: M1-10L
US-09-456-090A-46
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1 BIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA
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                                                                                                                                                                                            91.3%; Score 518.5; DB 2; Length 224; 94.4%; Pred. No. 6e-42; ive 2; Mismatches 3; Indels 1;
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APPLICANT: Buechler, Joe
APPLICANT: Balkirs, Gunars
APPLICANT: Balkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
FILE REFRENCE: 020015-000200US
CURRENT APPLICATION NUMBER: US/09/456,090A
CURRENT FILING DATE: 1999-12-06
NUMBER OF SEQ ID NOS: 110
SOCTWARE: Patentin Ver. 2.1
SEQ ID NO 88
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94.4%; Pred. No. 6e-42;
tive 2; Mismatches 3;
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 82
LENGTH: 224
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Best Local Similarity 94.4*
Matches 102; Conservative
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Matches 102; Conservative
                                                                                               ORGANISM: Homo sapiens
OTHER INFORMATION: M2-31L
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COTHER INFORMATION: M2-34L
US-09-456-090A-88
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US-09-456-090A-88
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US-09-456-090A-90
                                                                          TYPE: PRT
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APPLICANT: Huse, William
APPLICANT: Watkins, Jeffry
APPLICANT: Watkins, Jeffry
APPLICANT: Wu, Herren
TITLE OF INVENTION: Methods of Optimizing Antibody Variable Region Binding Affinity
FILE REFERENCE: AME-06352
CURRENT APPLICATION NUMBER: US/09/434,870
CURRENT FILING DATE: 2001-08-20
PRIOR APPLICATION NUMBER: 60/159,689
PRIOR APPLICATION NUMBER: 60/159,689
REGORD SEQ ID NOS: 4
SOFTWARE: Patentin Version 3.0
SEQ ID NO 2.
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                                                                                                                                                                                                                                                                            Length 224;
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Fatent No. 6680209
GENERAL INFORMATION:
APPLICANT: Buechler, Joe
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
TITLE OF INVENTION: HOMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
FILE REFREENCE: 020015-000200US
CURRENT APPLICANTION: 1999-12-06
CURRENT FILING DATE: 1999-12-06
NUMBER OF SEQ ID NOS: 110
                                                                                                                                                                                                                                                                       92.2%; Score 523.5; DB 2; Length 93.5%; Pred. No. 2e-42; tive 4; Mismatches 2; Indels
PRIOR APPLICATION NUMBER: US 60/157,415
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                    PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 46
LENGTH: 224
                                                                                                                                                                                                                                                                                                                           Matches 101; Conservative
                                                                                                                                                TYPE: PRT
OTHER INFORMATION: M1-10L
US-09-453-234-46
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Best Local Similarity
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US-09-434-870-2
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Length 224; Indels

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PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 88
LENGTH: 224
                                                                                                      TYPE: PRT
OCANISM: Homo sapiens
CTHER INFORMATION: M2-34L
US-09-453-234-88
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OTHER INFORMATION: M2-35L
US-09-453-234-90
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                                                                                                      91.3%; Score 518.5; DB 2; Length 224; 94.4%; Pred. No. 6e-42; Live 2; Mismatches 3; Indels 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     | JEAULI NO. 0.79132
| JEAULI NO. 0.79134
| APPLICANT: Buechler, Joe
| APPLICANT: Buchler, Gunars
| APPLICANT: Gray, Jeff
| APPLICANT: Lonberg, Nils
| APPLICANT: Geneharm International
| APPLICANT: Geneharm International
| TILE OF INVENTION: Human Antibodies
| FILE REFERENCE: 020015-0001100S
| CURRENT FILING DATE: 1999-12-01
| PRIOR APPLICATION NUMBER: US 60/157,415
| PRIOR PILING DATE: 1999-10-02
| NUMBER OF SEQ ID NOS: 112
| SEQ ID NO 82
| LENGTH: 224
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Patent No. 6794132
GENERAL INFORMATION:
APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Lonberg, Nils
APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: Grapharm International
TITLE OF INVENTION: Human Antibodies
FILE REFERENCE: 020015-0001100S
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                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 82, Application US/09453234 Patent No. 6794132
                                                                                                                           Best Local Similarity 94.4
Matches 102; Conservative
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; ORGANISM: Homo sapiens
; OTHER INFORMATION: M2-31L
US-09-453-234-82
  ; TYPE: PRT
; ORANISM: Homo sapiens
; OTHER INFORMATION: M2-35L
US-09-456-090A-90
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US-09-453-234-82
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US-09-453-234-88
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1 EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA 60
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Query Match 91.3%; Score 518.5; DB 2; Length 224; Best Local Similarity 94.4%; Pred. No. 6e-42; Matches 102; Conservative 2; Mismatches 3; Indels 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61 RPSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
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Patent No. 6680209
GENERAL INFORMATION:
APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Caray, Jeff
APPLICANT: Lonberg, Nils
APPLICANT: Conderg, Nils
APPLICANT: Conderg
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Pred. No. 6e-42;
2; Mismatches 3;
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APPLICANT: Buechler, Joe
APPLICANT: Buechler, Gunars
APPLICANT: Gray, Jdf;
APPLICANT: Lonberg, Nis
APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: Graph International
FILE REFERENCE: 020015-0001100S
CURRENT APPLICATION NUMBER: US/09/453,234
CURRENT APPLICATION NUMBER: US 60/157,415
PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SEQ ID NO 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 90, Application US/09453234
; Patent No. 6794132
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Best Local Similarity 94.4%;
Matches 102; Conservative 2
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Gaps

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1 BIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRAAGIPA 60
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                                                                                                                                                                                                                                                                                                  Query Match 90.6%; Score 514.5; DB 2; Length 224; Best Local Similarity 93.5%; Pred. No. 1.4e-41; Matches 101; Conservative 2; Mismatches 4; Indels 1;
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Batent No. 679432

GENERAL INFORMATION:
APPLICANT: Buechler, Joe
APPLICANT: Valkirs Gunars
APPLICANT: Canberg, Nils
APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: Genharm International
TITLE OF INVENTION: Human Antibodies
FILE REFERENCE: 020015-000110US
CURRENT APPLICATION NUMBER: US/09/453,234
CURRENT APPLICATION NUMBER: US/09/453,234
CURRENT APPLICATION NUMBER: US/09/453,234
FRIOR PILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SOFTWARE PATENTION OF SEQ ID NOS: 112
SEQ ID NOS 44

SEQ ID NO 84
FILE REFERENCE: 020015-000200US
CURRENT APPLICATION NUMBER: US/09/456,090A
CURRENT FILING DATE: 1999-12-06
NUMBER OF SEQ ID NOS: 110
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 84
LENGTH: 224
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Patent No. 6488934
PAPELICANT: HONG, Hyo Jeong
APPLICANT: FARK, Sung Sup
APPLICANT: KANG, Young Jun
APPLICANT: KANG, Young Jun
APPLICANT: KANG, Chang-Yuil
                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapiens
OTHER INFORMATION: M2-32L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Homo sapiens
OTHER INFORMATION: M2-32L
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                                                                                                                                                                                                                                                                                               Query Match 91.1%; Score 517.5; DB 2; Length 224; Best Local Similarity 94.4%; Pred. No. 7.5e-42; Matches 102; Conservative 1; Mismatches 4; Indels 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
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Patent No. 6680209
GENERAL INFORMATION:
APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Caray, Deff
APPLICANT: Lonberg, Nils
TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91.1%; Score 517.5; DB 2 94.4%; Pred. No. 7.5e-42; tive 1; Mismatches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Oray, Joff
APPLICANT: Oray, Joff
APPLICANT: General incommentational
APPLICANT: General incommentational
TITLE OF INVENTION: Human Antibodies
FILE REFERENCE: 020015-00011003
CURRENT APPLICATION NUMBER: US/09/453,234
CURRENT FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 36
LENGTH: 224
     CURRENT APPLICATION NUMBER: US/09/456,090A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 36, Application US/09453234
Patent No. 6794132
                         CURRENT FILING DATE: 1999-12-06
NUMBER OF SEQ ID NOS: 110
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 36
LENGTH: 224
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Best Local Similarity 94.4%
Matches 102; Conservative
                                                                                                                                                                                                                           ) OTHER INFORMATION: M1-1L
US-09-456-090A-36
                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Homo sapiens
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US-09-456-090A-84
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US-09-453-234-36
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US-09-453-234-36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 89.7%; Score 509.5; DB 2; Length 107;
Best Local Similarity 90.7%; Pred. No. 1.9e-41;
Matches 98; Conservative 5; Mismatches 4; Indels 1; Gaps
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Best Local Similarity 89.9%; Pred. No. 1.5e-39;
Matches 98; Conservative 3; Mismatches 7; Indels 1; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: Variable OTHER INFORMATION: region of light chain of human antibody (X82934)
APPLICANT: YOON, Sung Kwan

TITLE OF INVENTION: HUMANIZED ANTIBODY SPECIFIC FOR HUMAN 4-1BB AND

TITLE OF INVENTION: PHARMACEUTICAL COMPOSITION COMPRISING SAME
PILE REPERRINCE: 1303-124B

CURRENT APPLICATION NUMBER: US/09/438,954

CURRENT FILING DATE: 1999-11-12

SOFTWARE PILING DATE: 1999-11-12

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 40

LENGTH: 107

TYPE: PRT

ORGANISM: Artificial Sequence
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US-09-45-090A-38
US-09-45-090A-38
Sequence 38, Application US/09456090A
Sequence 38, Application US/09456090A
Sequence 38, Application US/09456090A
Sequence 38, Application US-09-6
Sequence 39, Application US-09-6
Septicant Description HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
CURRENT PILING DATE: 1999-12-06
SOFTWARE PECTATION NUMBER: US/09/456,090A
CURRENT FILING DATE: 1999-12-06
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 38
HURSH PATENT SECTION OF SECTION OF SEQ ID NO 38
HURSH PATENT SECTION OF 
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CRGANISM: Homo sapiens
CTHER INFORMATION: M1-3L
US-09-456-090A-38
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Appl Appl Appl Appli Appli

Sequence Sequence Sequence

Sequence Seq

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US-09-920-267C-8

Sequence 8, Application US/09920267C

Sequence 8, Application WS/09920267C

Publication No. US2003004004A1

GENERAL INFORMATION:

APPLICANT: Giles-Komar, Jill

APPLICANT: Trikha, Mohit

APPLICANT: Trikha, Mohit

APPLICANT: Nakada, Marian

TITLE OF INVENTION: ANTI-DUAL INTEGRIN ANTIBODIES, COMPOSITINS, METHODS AND USES

FILE REFERENCE: CEN 249

CURRENT APPLICATION NUMBER: US/09/920,267C

CURRENT APPLICATION NUMBER: 60/223,363

PRIOR PILING DATE: 2000-08-07

NUMBER OF SEQ ID NOS: 17

SOFTWARE: Patentin version 3.1

SERQ ID NO 8

TENERAL SECOND NOS: 17

SERQ ID NO 8
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Sequence 8, Application US/10305347A

Sequence 8, Application No. US20030143603A1

FUBLICANT: 0168-Komar, Jill

APPLICANT: Gles-Komar, Jill

APPLICANT: BETAIN SCALLON

TITLE OF INVENTION: ANTI-TNF ANTIBODIES, COMPOSITIONS, METHODS AND USES; CURRENT APPLICANTION NUMBER: US/10/305,347A

CURRENT FILING DATE: 2002-11-26

NUMBER OF SEQ ID NOS: 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
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Pred. No. 1.8e-41;
0; Mismatches 1; Indels
US-10-293-418-1049
US-10-881-688-91
US-11-0-886-3138-2
US-311-090-731-92
US-09-453-234-46
US-09-453-234-46
US-09-791-153A-71
US-10-443-466A-25
US-09-791-153A-71
US-09-791-153A-71
US-10-697-390-2
US-10-697-300-2
US-10-687-799-55
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99.1%;
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Best Local Similarity 99.1
Matches 107; Conservative
        TYPE: PRT
ORGANISM: Homo sapiens
        US-09-920-267C-8
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Sequence 1049, Ap
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                                                                                                                                                                                                                           US-09-920-137F-8
568
1 BIVLTQSPATLSLSPGERAT......QQRSNWPPFTFGPGTKVDIK 108
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(gn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

(gn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

(gn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

(gn2_6/ptodata/1/pubpaa/US108_PUBCOMB.pep:*

(gn2_6/ptodata/1/pubpaa/US108_PUBCOMB.pep:*

(gn2_6/ptodata/1/pubpaa/US108_PUBCOMB.pep:*
                         GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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US-10-362-347A-8

US-10-954-900A-8

US-11-009-731-90

US-10-408-901-24

US-10-656-769-12

US-10-656-769-12

US-10-656-769-13

US-10-656-769-13

US-10-408-901-8

US-10-408-901-8

US-10-408-901-8

US-10-408-901-16

US-11-102-403-11

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US-10-408-901-16

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US-09-880-748-1049
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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1 EIVLIQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA 60
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                                                                                                                                                                                                                                                                                                                                                                                      98.4%; Score 559; DB 5; Length 108; 99.1%; Pred. No. 1.8e-41; vative 0; Mismatches 1; Indels
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APPLICANT: PASSAWORE, David
APPLICANT: RANGAN, Vangipuram
APPLICANT: LANG, Thomas E.
APPLICANT: LANG, Thomas E.
APPLICANT: LANG, Michael T.
TITLE OF INVENTION: IP-10 ANTIBODIES AND THEIR USES FILE REFERENCE: MXI-312
CURRENT APPLICATION NUMBER: US/11/009,731
CURRENT FILING DATE: 2004-12-10
PRIOR APPLICATION NUMBER: 60/529180
PRIOR PILING DATE: 2003-12-10
NUMBER OF SEQ ID NOS: 126
SOFTWARE: PASSAGE OF Windows Version 4.0
SEQ ID NO 90
PRIOR FILING DATE: 2000-08-07
PRIOR APPLICATION NUMBER: 60/236,826
PRIOR FILING DATE: 2000-09-29
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PATENTIN Ver 3.1
SEQ ID NO 8
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SRINIVASAN, Mohan
CARDARELLI, JOSEPhine M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: DESHPANDE, Shirkant APPLICANT: HUANG, Haichun
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Best Local Similarity 99.13
Matches 107; Conservative
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CORGANISM: Homo sapiens
US-11-009-731-90
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US-10-954-900A-8
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US-10-408-901-24
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sequence 8, Application US/10720323
sequence 8, Application No. US20040185507A1
sequence 9, Application No. US20040185507A1
set Carbon No. US20040185507A1
set Carbon No. US20040185507A1
set Carbon No. US2004018507A1
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Sequence 8, Application US/10954900A

Publication No. US20050123541A1

GENERAL INFORMATION:

APPLICANT: Giles-Komar, Jill

APPLICANT: David Knight

APPLICANT: David Knight

APPLICANT: Bernie Scalus

TITLE OF INVENTION: ANTI-TYP ANTIBODIES, COMPOSITIONS, METHODS AND USES

FILE REFERENCE: CEN0250 DIV.S/10/954,900A

CURRENT APPLICATION NUMBER: US/10/954,900A

CURRENT FILING DATE: 2004-09-30

PRIOR APPLICATION UNMERE: 09/920,262

PRIOR APPLICATION NUMBER: 60/223,360
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                                                                                                                                                                                                               Query Match 98.4%; Score 559; DB 4; Length 100
Best Local Similarity 99.1%; Pred. No. 1.8e-41;
Matches 107; Conservative 0; Mismatches 1; Indels
                                                                            ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-305-347A-8
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CRGANISM: Homo sapiens
US-10-720-323-8
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US-10-720-323-8
                                    LENGTH: 108
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       SEQ ID NO 8
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Gaps

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Sequence 16, Application US/10394471B
; Sequence 16, Application US/10394471B
; Publication No. US20040185047A1
; Fublication No. US20040185047A1
; APPLICANT Glies-Komar, Jill; Carton, Jill; Scallon, Bernard J.
; TITLE OF INVENTION: ANTI-TNF ANTIBODIES, COMPOSITIONS, METHODS AND USES; FILE REFERENCE: CEN0202
; FILE REFERENCE: CEN0202
; CURRENT APPLICATION NUMBER: US/10/394,471B
; CURRENT FILING DATE: 2003-03-26
; RIOR APPLICATION NUMBER: 60/367,903
; RIOR APPLICATION NUMBER: 60/367,903
; RIOR APPLICATION NUMBER: 50/30-16
; SEQ ID NO 16
; SEQ ID NO 16
; LENGTH: 130
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     APPLICANT: Witte, Alison
APPLICANT: Vezina, Chris
APPLICANT: Wong, Lu Min
APPLICANT: Wong, Lu Min
APPLICANT: Olan, Xueming
TITLE OF INVENTION: Therapeutic Human Anti-IL-IR Monoclonal Antibody
FILE REFERENCE: 01,1554
CURRENT PPLLICANT: 2003-09-05
NUMBER OF SEQ ID NOS: 79
SOFTWARE: PatentIn version 3.0
SEQ ID NO 12
LENGTH: 128
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Publication No. US20040097712A1
GENERAL INFORMATION:
APPLICANT: Varian, Brian
APPLICANT: Witte, Alison
APPLICANT: Weakina, Chris
APPLICANT: Wong, Lu Min
APPLICANT: Qian, Xueming
TITLE OF INVENTION: Therapeutic Human Anti-IL-IR Monoclonal Antibody
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Pred. No. 9.8e-40;
1; Mismatches 3; Indels
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ilarity 96.3%;
Conservative 1
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; ORGANISM: Homo sapiens
US-10-656-769-12
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US-10-394-471B-16
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Best Local (
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APPLICANT: Elliot, Robin
APPLICANT: Sullivan, John
APPLICANT: Madlock, Eugene
APPLICANT: Madlock, Eugene
APPLICANT: Matin, Francis
TITLE OF INVENTION: Human Anti-OPGL Neutralizing Antibodies As Selective OPGL Pathway
TITLE OF INVENTION: Inhibitors
FILE REPERSION: Inhibitors
FILE REPERSION: MABER: US/10/408,901
CURRENT APPLICATION NUMBER: 108/10/408,901
NUMBER OF SEQ ID NOS: 76
SOFTWARE: Patentin version 3.0
SEQ ID NO 24
LENGRIH: 109
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APPLICANT: Hund, Haichun
APPLICANT: Hund, Haichun
APPLICANT: Bloot, Robin
APPLICANT: Bloot, Robin
APPLICANT: Sullivan, John
APPLICANT: Medlock, Eugene
APPLICANT: Medlock, Eugene
APPLICANT: Medlock, Eugene
APPLICANT: Medlock, Eugene
TITLE OF INVENTION: Inhibitors
FILE REPRENCE: MBHB 01-1145-A
CURRENT APPLICATION NUMBER: 2010/408,901
CURRENT APPLICATION NUMBER: 2003-04-07
NUMBER OF SEQ ID NOS: 76
SEQ ID NO 48
LENGTH: 215
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Pred. No. 6.7e-40;
3; Mismatches 2; Indels
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US-10-408-901-48
Sequence 48, Application US/10408901
Publication No. US20040023313A1
GENERAL INFORMATION
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; Sequence 12, Application US/10656769
; Publication No. US20040097712A1
; GENERAL INFORMATION:
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Best Local Similarity 95.4%;
Matches 103; Conservative
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CRGANISM: Homo sapiens
US-10-408-901-24
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; ORGANISM: Homo sapiens
US-10-408-901-48
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                                                                                                                                                                                                                                                                                                                                                                                           Score 539; DB 4; Length 215;
Pred. No. 2e-39;
4; Mismatches 2; Indels
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APPLICANT: Lu, Chafen; TITLE OF INVENTION: Antibodies that bind alphaE Integrin; FILE REFERENCE: 1855.2025-000; CURRENT APPLICATION NUMBER: US/10/173,551; CURRENT FILING DATE: 2002-06-14; NUMBER OF SEQ ID NOS: 70; SOFTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 9.
                    PILE REFERENCE: MHHB 01-1145-A
CURRENT APPLICATION NUMBER: US/10/408,901
CURRENT FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 76
SOFTWARE: Patentin version 3.0
SEQ ID NO 32
LENGTH: 215
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TITLE OF INVENTION: Inhibitors
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Best Local Similarity 94.4%;
Matches 102; Conservative 4
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Best Local Similarity 96.34
Matches 104; Conservative
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NAME/KEY: SITE
LOCATION: (24)...(34)
OTHER INFORMATION: CDR1
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LOCATION: (50)...(56)
OTHER INFORMATION: CDR2
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; OTHER INFORMATION: CDR3
US-10-173-551-9
                                                                                                                                                                                                                                                                   TYPE: PRT
CORGANISM: Homo sapiens
US-10-408-901-32
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ORGANISM: Homo sapiens
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NAME/KEY: SITE
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US-10-886-838-4
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Publication No. US20040023313A1
GENERAL INFORMATION:
APPLICANT: BOyle, William
APPLICANT: Elliot, Robin
APPLICANT: Medick, Eugene
APPLICANT: Medick, Eugene
APPLICANT: Medick, Eugene
APPLICANT: Martin, Francis
TITLE OF INVENTION: Human Anti-OPGL Neutralizing Antibodies As Selective OPGL Pathway
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APPLICANT: BOyle, William

APPLICANT: Huang, Haichun

APPLICANT: Bliot, Robin

APPLICANT: Bliot, Robin

APPLICANT: Sullivan, John

APPLICANT: Medlock, Eugene

APPLICANT: Medlock
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                                                                                                                                                                                                                                                                                                                                                                       Length 235;
                                                                                                                                                                                                                                                                                                                                                                                                                                        3; Indels
                                                                                                                                                                                                                                                                                                                                                                   95.1%; Score 540; DB 4;
96.3%; Pred. No. 1.7e-39;
trive 1; Mismatches 3.
                              CURRENT APPLICATION NUMBER: US/10/656,769
CURRENT FILING DATE: 2003-09-05
NUMBER OF SEQ ID NOS: 79
SOFTWARE: PATENTIA NOS: 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 8, Application US/10408901
Publication No. US20040023313A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 94.4
Matches 102; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 104; Conservative
                                                                                                                                                                                                                                 TYPE: PRT
CORGANISM: Homo sapiens
US-10-656-769-38
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ORGANISM: Homo sapiens
   FILE REFERENCE: 01,1554
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 12
US-10-408-901-32
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                                                                                                                                                                SEQ ID NO 38
LENGTH: 235
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US-10-408-901-8
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APPLICANT: Hoffmann-La Roche Inc.

TITLE OF INVENTION: Antibodies against insulin-like growth factor I receptor and uses FITLE OF INVENTION: thereof
FILE OP INVENTION: thereof
FILE SPERENCE: 21695
CURRENT APPLICATION NUMBER: US/10/886,838
CURRENT FILING DATE: 2004-07-08
PRIOR APPLICATION NUMBER: BP 03015526
PRIOR PILING DATE: 2003-07-10
NUMBER OF SEQ ID NOS: 8
SEQ ID NO 4
LENGTH: 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: COMPOSITIONS, METHODS AND KITS RELATING TO ANTI-PLATELET
TITLE OF INVENTION: COMPOSITIONS, METHODS AND KITS RELATING TO ANTI-PLATELET
TITLE OF INVENTION: AUTOANTIBODIES AND INHIBITORS THEREOF
TITLE OF INVENTION: AUTOANTIBODIES AND INHIBITORS THEREOF
TITLE OF INVENTION: 033893-5050
CURRENT PRILING DATE: 2004-12-23
FRIOR PILING DATE: 2003-07-03
FRIOR FILING DATE: 2003-07-03
FRIOR PILING DATE: 2002-09-18
FRIOR PILING DATE: 2002-09-18
FRIOR PILING DATE: 2002-09-18
FRIOR PILING DATE: 2002-09-18
FRIOR FILING DATE: 2002-09-18
FRIOR PILING DATE: 2003-09-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 BIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 94.4%; Score 536; DB 5; Length 10: Best Local Similarity 95.4%; Pred. No. 1.8e-39; Matches 103; Conservative 2; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-021-715-101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapiens
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US-11-021-715-101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: HC
US-10-886-838-4
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Sequence Sequence Sequence

Sequence 178, Appli Sequence 178, Appli Sequence 1534, Appli Sequence 165, Appl Sequence 127, Appl Sequence 177, Appl Sequence 177, Appl Sequence 177, Appl Sequence 18, Appli Sequence 88, Appli Sequence 88, Appl Sequence 135, Appl Sequence 135, Appl Sequence 135, Appl Sequence 135, Appl Sequence 115, Appl Sequence 115, Appl Sequence 115, Appl Sequence 116, Appl Sequence 161, Appl Sequence 1

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Sequence:

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RPSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFFGPGTKVDIK 108
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; Sequence 23, Application US/11093274
; Publication No. US23050266008A1
; GENERAL INFORMATION:
; APPLICANT: Graziano, Robert
; APPLICANT: Cardarelli, Josephine M.
; APPLICANT: Cutter, Beth
; APPLICANT: Cutter, Beth
; APPLICANT: Stinivasan, Mohan
; TITLE OF INVENTION: IRTA-5 ANTIBODIES AND THEIR USES
; FILE REFERENCE: 04280/1201101-US1
; CURRENT FILING DATE: 2005-03-28
; PRIOR PILING DATE: 2005-03-29
; PRIOR PILING DATE: 2004-03-29
; NUMBER OF SEQ ID NOS: 41
; SOFTWARE: Patentin version 3.2
; SEQ ID NO 23
; LENGTH: 108
; TVDP: DDP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 94.9%; Score 539; DB 7; L. Similarity 95.4%; Pred. No. 4.6e-35; 33; Conservative 3; Mismatches 2;
US-11-054-515-1264
US-11-064-515-1219
US-11-064-174-189
US-11-074-186-7
US-11-074-186-7
US-11-054-515-905
US-11-054-669-84
US-11-054-669-85
US-11-084-554-127
US-11-084-554-137
US-11-084-554-137
US-11-074-669-88
US-11-072-320-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: KIRIN BEER KABUSHIKI KAISHA
TITLE OF INVENTION: ANTI TRAIL-R ANTIBODY
FILE REFERENCE: PH-1573-PCT
CURRENT PEPLICATION NUMBER: US/10/721,763
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: JP2001-150213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19, Application US/10721763
Publication No. US20050249729A1
GENERAL INFORMATION:
        ; ORGANISM: Homo sapiens
US-11-093-274-23
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Best Local S:
Matches 103
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Sequence 805, App
Sequence 806, App
Sequence 807, App
Sequence 8, Appli
Sequence 1, Appli
Sequence 1268, Appli
Sequence 1268, Appli
                                                                                                                                                                                     ; Search time 6.92308 Seconds (without alignments) 116.825 Million cell updates/sec
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                                                                                                                                                                                                                                                                                           US-09-920-137F-8
568
1 BIVLTQSPATLSLSPGERAT......QQRSNWPPFTFGPGTKVDIK 108
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1: /cgn2 6/ptodata/1/pubpaa/USO6 NEW PUB.pep:*
2: /cgn2 6/ptodata/1/pubpaa/USO6 NEW PUB.pep:*
3: /cgn2 6/ptodata/1/pubpaa/USO7 NEW PUB.pep:*
4: /cgn2 6/ptodata/1/pubpaa/PCT NEW PUB.pep:*
5: /cgn2 6/ptodata/1/pubpaa/USO9 NEW PUB.pep:*
6: /cgn2 6/ptodata/1/pubpaa/USO9 NEW PUB.pep:*
7: /cgn2 6/ptodata/1/pubpaa/USO1 NEW PUB.pep:*
8: /cgn2 6/ptodata/1/pubpaa/USO1 NEW PUB.pep:*
                                   GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd
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US-10-721-763-19

US-11-054-215-1049

US-11-093-274-22

US-10-721-763-23

US-10-721-763-23

US-11-040-159-17

US-11-040-159-17

US-11-040-159-17

US-11-044-569-86

US-11-054-569-87

US-11-054-669-87

US-11-054-669-87

US-11-054-669-87

US-11-054-669-87

US-11-054-669-87

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US-11-054-515-1952
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   57103 seqs, 7488799 residues
                                                                                                                                                                                           December 28, 2005, 14:04:11
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Minimum DB Maximum DB

Database

Result Š

Searched:

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Gaps

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Length 108; Indels

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; ORGANISM: Homo sapiens
US-11-093-274-22
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US-11-054-515-1049
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US-11-093-274-22
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; Publication No. US2005025532A1
; GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
; FILE REFERENCE: PF533P3
; CURRENT APPLICATION NUMBER: US/11/054,515
; CURRENT FILING DATE: 2005-02-10
                                                                                                                                                                                                                                                                                                                                                                                                                                 61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                        81 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNW-PLTFGPGTKVDIK 127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
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                                                                                                                                                                                                                                                                                                              2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 24, Application US/11093274

| Sequence 24, Application US/11093274
| Publication No. US200506008A1
| GENERAL INFORMATION:
| APPLICANT: Cardarelli, Josephine M. APPLICANT: Cardarelli, Josephine M. APPLICANT: Cutter, Beth
| APPLICANT: Srinivasan, Mohan | TILE OF INVENTION: IRRA-5 ANTIBODIES AND THEIR USES FILE REFERENCE: 04280/1201101-US1
| CURRENT APPLICATION NUMBER: US/11/093,274
| CURRENT PILING DATE: 2005-03-28
| PRIOR FILING DATE: 2004-03-29
| WIMBER OF SEQ ID NOS: 41
| SOFTWARE: PatentIn version 3.2
| SEQ ID NO 24
                                                                                                                                                                                                                                                                    93.6%; Score 531.5; DB 6 96.3%; Pred. No. 1.9e-34; iive 1; Mismatches 2
; PRIOR FILING DATE: 2001-05-18
; PRIOR APPLICATION NUMBER: JP2001-243040
; PRIOR FILING DATE: 2001-08-09
; PRIOR FILING DATE: 2001-10-11
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 19
; LENGTH: 129
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Best Local Similarity 96.34
Matches 104; Conservative
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                                                                                                                                                                                  TYPE: PRT; ORGANISM: Homo sapiens
US-10-721-763-19
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; ORGANISM: Homo sapiens
US-11-093-274-24
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US-11-054-515-1049
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PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
PRIOR PLILING DATE: 2004-02-11
PRIOR PAPLICATION NUMBER: 60/580,347
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR PILING DATE: 2001-11-16
PRIOR PLING DATE: 2001-12-19
PRIOR PLING DATE: 2001-12-19
PRIOR PLING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR PLING DATE: 2001-06-25
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR PLING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR PLING DATE: 2001-03-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61 RPSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93.1%; Score 529; DB 7; Length 251; 95.4%; Pred. No. 5e-34;
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Sequence 2.7, 4.4.4.

Publication No. US20050266008A1

SEGRERAL INFORMATION:

APPLICANT: Gardarelli, Josephine M.

APPLICANT: Kempe, Thomas

APPLICANT: Cutter, Beth

APPLICANT: Cutter, Beth

TITLE OF INVENTION: IRTA-5 ANTIBODIES AND THEIR USES

FILE REFERENCE: 04280/1201101-US1

CURRENT APPLICATION NUMBER: 0557,741

PRIOR APPLICATION NUMBER: 60/557,741

PRIOR PLING DATE: 2004-03-29

NUMBER OF SEQ ID NOS: 41

SEQ ID NO 22

LENGTH: 108
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Best Local Similarity 95.4<sup>1</sup>
Matches 103; Conservative
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; ORGANISM: Homo sapiens
US-11-054-669-86
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Score 522.5; DB 6; Length 128;
Pred. No. 9.1e-34;
1; Mismatches 3; Indels 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      81 RPSGSGSGTDFTLTISSLEPEDEPAVYYCQQRSNW-PLTFGGGTKVEIK 127
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Sequence 23, Application US/10721763

Sequence 23, Application US/10721763

Publication No. US20050249729A1

GENERAL INFORMATION:
APPLICANT: KIRIN BEER KABUSHIKI KAISHA

TITLE NOF INVENTION: ANTI TRAIL-R ANTIBODY
FILE REFERENCE: PH-1573-PCT
CURRENT APPLICATION NUMBER: US/10/721,763

CURRENT FILING DATE: 2003-11-26

PRIOR PILING DATE: 2001-05-18

PRIOR PILING DATE: 2001-05-18

PRIOR FILING DATE: 2001-06-09

PRIOR FILING DATE: 2001-08-09

PRIOR FILING DATE: 2001-08-09

PRIOR FILING DATE: 2001-10-11

NUMBER OF SEQ ID NOS: 45

SEQ ID NO 23
                                                                                                                                                                                          TITLE OF INVENTION:
TITLE OF INVENTION: ANTI TRAIL-R ANTIBODY
FILE REPERENCE: PH-1573-PCT
CURRENT APPLICATION NUMBER: US/10/721,763
CURRENT FILING DATE: 2003-11-26
PRIOR PRIOR APPLICATION NUMBER: JP2001-150213
PRIOR PLING DATE: 2001-05-18
PRIOR PLING DATE: 2001-06-09
PRIOR PLING DATE: 2001-08-09
PRIOR PLING DATE: 2001-08-09
PRIOR PLING DATE: 2001-01
NUMBER: OF SEQ ID NOS: 45
SEQ ID NO 31
                                                                                                                                 ; Sequence 31, Application US/10721763; Publication No. US20050249729A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92.0%;
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Matches 102; Conservative
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Best Local Similarity 95.4
Matches 103; Conservative
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; ORGANISM: Homo sapiens
US-10-721-763-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Homo sapiens
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RESULT 8

RESULT 9

1 SEQUENCE 17, APPLICATION SILENDEDPAYTYCORSNAPPTFORTKUIK 127

1 SEQUENCE 17, APPLICATION NO. US2005025552A1

1 SERVICANT SHALL SCHOOL STATE AND STATE APPLICANT SECRETARIAN SHALLOWS SECRET
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US-11-093-274-33
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                                                                                                               1 EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA 60
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                                                                                        1 BIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA 60
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                                               0; Gaps
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is Sequence 138, Application US/11084554

sequence 138, Application US/11084554

spublication No. US20050266079A1

GENERAL INFORMATION:

APPLICANT: Reliermann, Sirid-Ai

APPLICANT: Green, Larry L.

APPLICANT: Green, Larry L.

APPLICANT: Korver, Wouter

ITILE OF INVENTION: REDUCING THE RISK OF HUMAN ANTI-HUMAN

TITLE OF INVENTION: AAVIBODIES THROUGH V GENE MANIPULATION

FILE REPERENCE: ABGENIX.100A

CURRENT PELLING DATE: 2005-03-17

PRIOR APPLICATION NUMBER: 60/554,372

PRIOR APPLICATION NUMBER: 60/554,372

PRIOR APPLICATION NUMBER: 60/574,661

PRIOR APPLICATION NUMBER: 60/574,661

PROPER PRIOR PELING DATE: 2004-03-19

ROPTWARE OF SEQ ID NOS: 266

SEQ ID NO 138

LENGTH: 95
Score 487; DB 7; Length 95;
Pred. No. 3.4e-31;
0; Mismatches 1; Indels
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                                                                                                                                                                                 61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWP 95
                                                                                                                                                                                                     61 RFSGSGGTDFTLTISSLEPEDFAVYYCQQRSNWP 95
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APPLICANT: Graziano, Robert
APPLICANT: Graziano, Robert
APPLICANT: Cardarelli, Josephine M.
APPLICANT: Cutter, Beth
APPLICANT: Cutter, Beth
TITLE OF INVENTION: IRRA-5 ANTIBODIES AND THE
TITLE REPREBNCE: 04280/120110.143.
CURRENT FILING DATE: 2005-03-28
FRIOR PLICATION NUMBER: 05/557,741
PRIOR PLILING DATE: 2004-03-29
NUMBER OF SEQ ID NOS: 41
SOFTWARE: Patentin version 3.2
SENGTHARE: 94
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Query Match 85.7%;
Best Local Similarity 98.9%;
Matches 94; Conservative (
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Best Local Similarity 98.9
Matches 94; Conservative
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US-11-084-554-138
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ORGANISM: Homo sapiens
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1 BIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA 60
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| Sequence 131, Application US/11084554
| Publication No. US20050260679A1
| GENERAL INFORMATION:
| APPLICANT: Kellermann, Sirid-Ai
| APPLICANT: Korver, Wouter
| TILLE OF INVENTION: REDUCING THE RISK OF HUMAN ANTI-HUMAN TITLE OF INVENTION: REDUCING THE RISK OF HUMAN ANTI-HUMAN TITLE OF INVENTION: MURBER: US/11/084,554
| CURRENT APPLICATION NUMBER: US/11/084,554
| CURRENT FILING DATE: 2004-03-19
| PRIOR PAPLICATION NUMBER: 60/554,372
| PRIOR PAPLICATION NUMBER: 60/554,661
| PRIOR PILING DATE: 2004-03-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   82.9%; Score 471; DB 7; Length 95; 96.8%; Pred. No. 5.5e-30; ive 0; Mismatches 3; Indels
       Length 94;
                                                                       1; Indels
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FUblication No. US20050261480A1
GURERAL INFORMATION:
APPLICANT: FOOCE, Jefferson
TITLE OF INVENTION: SUPER HUMANIZED ANTIBODIES
FILE REFERENCE: 30219/US/3
CURRENT FILING DATE: 2005-02-08
FRIOR APPLICATION NUMBER: US 10/194,975
FRIOR APPLICATION NUMBER: US 60/305,111
FRIOR APPLICATION NUMBER: US 60/305,111
FRIOR APPLICATION NUMBER: US 60/305,111
FRIOR PILING DATE: 2001-07-12
FRIOR FILING DATE: 2001-07-12
SUPUMBER OF SEQ ID NOS: 124
SEQ ID NO 87
LENGTH: 95
Score 480; DB 7;
Pred. No. 1.1e-30;
0; Mismatches 1.
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   Query Match
Best Local Similarity 98.9%;
Matches 93; Conservative
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Matches 91; Conservative
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US-11-054-669-87
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ORGANISM: Homo sapiens
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US-11-000-463-332
; Sequence 33.2, Application US/11000463
; Publication No. US20050266423A1
; GENERAL INFORMATION:
    APPLICANT: Tang, Y Tom
    APPLICANT: Liu, Chenghua
; APPLICANT: Chenghua
; APPLICANT: Gran, Rui-hong
; APPLICANT: Qian, Xiaohong B.
APPLICANT: Wehrman, Tom
; APPLICANT: Zhang, Jie
; APPLICANT: Zhang, Jie
; APPLICANT: Zhou, Ping
; APPLICANT: Zhou, Ping
; APPLICANT: Zhou, Ping
; APPLICANT: Dramanac, Radoje T.
; TTLE OF INVENTION: Novel Nucleic Acids and Polypeptides
; TILE REFERENCE: 785CIPP4CN
                                          Query Match 82.9%; Score 471; DB 7; Length 95; Best Local Similarity 96.8%; Pred. No. 5.5e-30; Matches 91; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                        61 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNW 94
                                                                                                                                                                                                                                                                        61 RFSGSGPGTDFTLTISSLEPEDFAVYYCQQRSNW 94
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Publication No. US20050249729A1

GENERAL INFORMATION:

APPLICANT: KIRIN BEER KABUSHIKI KAISHA

TITLE OF INVENTION: ANTI TAALL-R ANTIBODY

FILE REFERENCE: PH-1573-PCT

CURRENT APPLICATION NUMBER: US/10/721,763

CURRENT FILING DATE: 2001-05-18

FRIOR FILING DATE: 2001-05-18

FRIOR FILING DATE: 2001-08-09

FRIOR FILING DATE: 2001-08-09

FRIOR FILING DATE: 2001-08-09

FRIOR FILING DATE: 2001-08-09

FRIOR FILING DATE: 2001-10-11

NUMBER OF SEQ ID NOS: 45

SEQ ID NO 27

SEQ ID NO 27
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CURRENT FILING DATE: 2004-11-29
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PRIOR APPLICATION NUMBER: 10/291,265
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ORGANISM: Homo sapiens
US-11-084-554-131
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US-10-721-763-27
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Published_Applications Mucleic Acid and Published_Applications Amino Acid database searches now generate two sets of results each. The Published_Applications databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Newly published applications will appear in the Published Applications May databases; older published

Searches run against Aucleic, Acid Published Applications produce two sets of results, with the extensions trapbm (Published Applications NA Mew).

Searches run against Amino Acid Published Applications produce two sets of results, with the extensions Searches run against Amino Acid Published Applications produce two sets of results, with the extensions are produced and applications AA Mew).

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Sequence 13. Appl. Sequence 15. Appl. Sequence 96. Appl. Sequence 18. Appl. Sequence 111. Appl. Sequence 111. Appl. Sequence 111. Appl. Sequence 1110. Appl. Sequence 1110. Appl. Sequence 1110. Appl. Sequence 1519. Appl. Sequence 956. Appl. Sequen

Sequence Seq

1109, Ap 1421, Ap 5, Appli

Sequence Sequence 1 Seguence

Sequence:

Run on:

Searched:

Database

Result)

No.

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ADSVKGRFTISRDNSKYTLYLØMNSLRAEDTAVYYCARDR----GIAAGGNYYYYGMDVW 116
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:

JETUE ADELICANT: Ruben et al.

TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS

FILE REPRENEUR: PES2383

CURRENT APPLICATION NUMBER: US/11/054,515

CURRENT FILING DATE: 2005-02-10

PRIOR PILING DATE: 2004-06-18

PRIOR PILING DATE: 2004-06-18

PRIOR PILING DATE: 2004-06-18

PRIOR PILING DATE: 2001-11-4

PRIOR PILING DATE: 2001-11-4

PRIOR PILING DATE: 2001-11-4

PRIOR PILING DATE: 2001-11-14

PRIOR PILING DATE: 2001-11-14

PRIOR PILING DATE: 2001-11-15

PRIOR PILING DATE: 2001-06-15

PRIOR PILING DATE: 2001-06-15

PRIOR PILING DATE: 2001-06-15

PRIOR PILING DATE: 2001-06-15

PRIOR PILING DATE: 2001-03-21

PRIOR PILING DATE: 2001-03-21

PRIOR PILING DATE: 2001-03-21

PRIOR PILING DATE: 2001-03-16

PRIOR PILING DATE: 2001-03-16
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SEQ ID NO 1731
LENGTH: 252
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llarity 86.9%; Pred. No. 5.5e-42;
Conservative 2; Mismatches 10;
US-11-054-515-1105
US-11-054-515-1108
US-11-064-174-13
US-11-064-174-152
US-11-1054-515-896
US-11-155-775-18
US-11-155-775-18
US-11-064-174-146
US-11-054-174-146
US-11-054-174-146
US-11-054-174-174
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US-11-054-175-181
US-11-054-515-956
US-11-054-515-881
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US-11-054-515-881
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Publication No. US20050255532A1
GENERAL INFORMATION:
   : Homo sapiens
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Best Local Similarity
Matches 113; Conserv
                 US-11-054-515-1731
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330, Ap
839, Ap
1394, Ap
981, App
140, App
8, Appli
397, App
1115, Ap
1724, Ap
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                                                                                                                 December 28, 2005, 14:04:11; Search time 8.07692 Seconds (without alignments) 116.825 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpaa/USO8 NEW PUB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/USO8 NEW PUB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/USO7 NEW PUB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/NESO NEW PUB.pep:*
7: /cgn2_6/ptodata/1/pubpaa/USIO NEW PUB.pep:*
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8: /cgn2_6/ptodata/1/pubpaa/USIO NEW_PUB.pep:*
8: /cgn2_6/ptodata/1/pubpaa/USIO NEW_PUB.pep:*
                   GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd
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Maximum Match 100%
Listing first 45 summaries
                                                                                   - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                            seq length: 0
seq length: 200000000
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Match 1
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Perfect score:
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Maximum DB
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TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

FILE REFERENCE: 09596-4202

CURRENT APPLICATION NUMBER: US/11/064,174

CURRENT FILING DATE: 2005-02-22

PRIOR APPLICATION NUMBER: US/09/240,274

PRIOR APPLICATION NUMBER: 60/091,380

PRIOR FILING DATE: 1999-01-29

PRIOR *PLING DATE: 1999-01-10

PRIOR *APPLICATION NUMBER: 60/028,550

PRIOR *APPLICATION NUMBER: 60/028,550

PRIOR *PLING DATE: 1996-10-11

NUMBER OF SEQ ID NOS: 224

SOFTWARE: PALENTIN VEY: 2.0

LENGTH: 126
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Sequence 512, Application US/11054515

Publication No. US2005025532A1

GENERAL INFORMATION:

APPLICAMT: Ruben et al.

TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS

FILE REPERENCE: PF523P3

CURRENT APPLICATION NUMBER: US/11/054,515

CURRENT APPLICATION NUMBER: G0/543,296

PRIOR APPLICATION NUMBER: 60/580,347

PRIOR FILING DATE: 2004-06-18

PRIOR FILING DATE: 2002-11-14

PRIOR FILING DATE: 2002-11-14

PRIOR APPLICATION NUMBER: 60/340,817

PRIOR FILING DATE: 2001-11-16

PRIOR APPLICATION NUMBER: 60/340,817

PRIOR APPLICATION NUMBER: 60/340,817

PRIOR FILING DATE: 2001-12-19

PRIOR FILING DATE: 2001-12-19

PRIOR FILING DATE: 2001-12-19

PRIOR FILING DATE: 2001-12-19

PRIOR FILING DATE: 2001-06-15
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Best Local Similarity 84.1%; Pred. No. 1.2e-41;
Matches 106; Conservative 8; Mismatches 12; Indels
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US-11-064-174-26
                                                                                                                                                                                                        ; Sequence 26, Application US/11064174; Publication No. US20050282252A1; GENERAL INFORMATION:
                                               117 GOGTTVTVSS 126
                                                                          |:|| |||||
120 GRGTLVTVSS 129
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ORGANISM: Homo sapiens
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US-11-064-174-26
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GENERAL IN CACHAIN.

APPLICANT: Siegel, Donald L.

TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

FILE REFERENCE: 09596-4202

CURRENT APPLICATION NUMBER: US/11/064,174

CURRENT APPLICATION NUMBER: US/09/240,274

PRIOR APPLICATION NUMBER: 60/081,380

PRIOR FILING DATE: 1999-01-29

PRIOR FILING DATE: 1998-04-10

PRIOR FILING DATE: 1996-10-11

NUMBER OF SEQ ID NOS: 224

SOFTWARE: PATENTIN Ver. 2.0

LENGTH: 126
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NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 512
LENGTH: 249
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                                                                                                                                                                                                                                                                                                                                                                                        Length 249;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              , OTHER INFORMATION: anti-Rh(D) antibody clone SH56 US-11-064-174-153
                                                                                                                                                                                                                                                                                                                                                                                     Score 560; DB 7;
Pred. No. 6.1e-41;
2; Mismatches 15;
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR PILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
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ilarity 86.5%;
Conservative 2
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US-11-054-515-512
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Best Local Similarity
Matches 109; Conserv
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Sequence 839, Application US/11054515; Publication No. US20050255332A1; GENERAL INFORMATION:
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84.5%;
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Best Local Similarity 84.5<sup>1</sup>
Matches 109; Conservative
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                                                                                                                                                                                                                ; ORGANISM: Homo sapiens
US-11-054-515-1330
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US-11-054-515-839
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Best Local Similarity
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                                                                                                          Sequence 25, Application US/11064174

Sequence 25, Application US/11064174

Publication No. US2005028252A1

GENERAL INFORMATION:

TITLE OF INVENTION: SCRTING METHOD FOR PRODUCTION THEREOF

TITLE OF INVENTION: SCRTING METHOD FOR PRODUCTION THEREOF

TITLE OF INVENTION: MUMBER: US/11/064,174

CURRENT FILING DATE: 2005-02-22

PRIOR PAPLICATION NUMBER: US/09/240,274

PRIOR PAPLICATION NUMBER: 60/081,380

PRIOR APPLICATION NUMBER: 60/081,380

PRIOR APPLICATION NUMBER: 60/081,380

PRIOR APPLICATION NUMBER: 60/028,550

PRIOR APPLICATION NUMBER: 60/028,550

PRIOR APPLICATION NUMBER: 60/028,550

PRIOR APPLICATION NUMBER: 60/028,550

SEQ ID NOS: 224

SEQ ID NO 25

LENGTH: 126
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GENERAL INFORMATION:

TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS

FILE REFERENCE: PF533P3

CURRENT APPLICATION NUMBER: US/11/054,515

CURRENT FILING DATE: 2005-02-10

FRIOR APPLICATION NUMBER: 60/543,296

FRIOR FILING DATE: 2004-02-11

FRIOR APPLICATION NUMBER: 60/580,347

FRIOR FILING DATE: 2004-06-18

FRIOR FILING DATE: 2002-11-14

FRIOR FILING DATE: 2001-11-14

FRIOR FILING DATE: 2001-11-16

FRIOR FILING DATE: 2001-11-16

FRIOR FILING DATE: 2001-11-219

FRIOR FILING DATE: 2001-11-219

FRIOR FILING DATE: 2001-06-15

FRIOR FILING DATE: 2001-06-15

FRIOR FILING DATE: 2001-06-15

FRIOR FILING DATE: 2001-06-25

FRIOR FILING DATE: 2001-06-25

FRIOR FILING DATE: 2001-06-25
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Pred. No. 5.8e-41;
8; Mismatches 13; Indels
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US-11-064-174-25
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Best Local Similarity 83.3%;
Matches 105; Conservative 8
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ORGANISM: Homo sapiens
121 TVTVSS 126
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61 ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARD-RGIAAGGNYYYYGMDVWGQG 119
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JITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REPRENCE: PF523P3
CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT PILING DATE: 2005-02-10
FRIOR PELICATION NUMBER: 60/543,296
FRIOR PELING DATE: 2004-02-11
FRIOR PELING DATE: 2004-02-11
FRIOR PELING DATE: 2004-02-11
FRIOR PELING DATE: 2004-02-11
FRIOR FILING DATE: 2001-03-14
FRIOR FILING DATE: 2001-11-16
FRIOR FILING DATE: 2001-11-16
FRIOR PELING DATE: 2001-10-16
FRIOR PELING DATE: 2001-06-25
FRIOR PELING DATE: 2001-06-25
FRIOR PELING DATE: 2001-06-25
FRIOR FILING DATE: 2001-03-21
FRIOR FILING DATE: 2001-03-16
FRIOR FILING DATE: 2001-03-21
FRIOR FILING DATE: 2001-03-21
FRIOR FILING DATE: 2001-03-16
FRIOR FILING DATE: 2001-03-17
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PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2000-10-17
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 3247
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Pred. No. 1.9e-40;
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Pred. No. 2e-40;
3; Mismatches
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                                                                                                                    DSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDR----GIAAGGNYYYYGMDVWG 117
                                                                                                                                                                DSVKGRPTISRDNSKNTLYLQMNSLRAEDTAVYYCARDREAYYDILTGYYLYYYYMDVWG 121
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| Sequence | 1394 | Application US/11054515 |
| Publication No. US2005025532A1 |
| GENERAL INFORMATION: |
| GENERAL INFORMATION: |
| APPLICAMT: Ruben et al. |
| TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS |
| TITLE OF INVENTION NUMBER: US/11/054,515 |
| CURRENT PILING DATE: 2005-02-10 |
| PRIOR PILING DATE: 2004-06-18 |
| PRIOR PELICATION NUMBER: 60/543,296 |
| PRIOR PILING DATE: 2004-06-18 |
| PRIOR PILING DATE: 2004-06-18 |
| PRIOR FILING DATE: 2001-11-14 |
| PRIOR PILING DATE: 2001-11-14 |
| PRIOR PILING DATE: 2001-11-14 |
| PRIOR PILING DATE: 2001-11-15 |
| PRIOR PILING DATE: 2001-12-19 |
| PRIOR PILING DATE: 2001-05-25 |
| PRIOR PILING DATE: 2001-05-25 |
| PRIOR PILING DATE: 2001-06-15 |
| PRIOR PILING DATE: 2001-06-15 |
| PRIOR PILING DATE: 2001-06-15 |
| PRIOR PILING DATE: 2001-03-21 |
| PRIOR PILING DATE: 2001-03-31 |
| PRIOR PILING DATE: 2001-03-31 |
| PRIOR PILING DATE: 2001-03-31 |
| PRIOR PILING DATE: 2001-03-16 |
| PRIOR PILING DATE: 2001-01-11 |
| PRIOR PILING DATE: 2001-03-16 |
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| PRIOR PILING DATE: 2001-01-11 |
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| PRIOR PILING DATE: 2001-03-16 |
| PRIOR PILING DATE: 2001-01-11 |
| PRIOR PIL
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SEQ ID NO 1394
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122 RGTTVTVSS 130
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KGTLVTVSS 129
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; ORGANISM: Homo sapiens
US-11-054-515-1394
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RESULT 9
US-11-054-515-981
Sequence 981, Application US/11054515
; Publication No. US2005025532A1
; GENERAL INFORMATION:
; APPLICANT: Ruben et al.

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Sequence 140, Application US/11064174

Publication No. US20050282521

GENERAL INFORMATION:
APPLICANT: Siegel, Donald L.
TITLE OF INVENTION:
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
THE REPRENCE TO 1995-02-22
PRIOR FILING DATE: 1999-01-29
PRIOR FILING DATE: 1999-04-10
PRIOR FILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224

NUMBER OF SEQ ID NOS: 224
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                   FILE OF INVENTION: ANTIDODISE CARC IMMUNOSPECIFICALLY BING BLYS
FILE REFRENCE: PF52193
CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT FILING DATE: 2005-02-10
PRIOR PLING DATE: 2004-02-11
PRIOR PLING DATE: 2004-02-11
PRIOR FILING DATE: 2004-06-18
PRIOR FILING DATE: 2004-06-18
PRIOR FILING DATE: 2001-06-18
PRIOR FILING DATE: 2001-11-16
PRIOR PLING DATE: 2001-11-16
PRIOR PLING DATE: 2001-11-16
PRIOR PLING DATE: 2001-11-19
PRIOR PLING DATE: 2001-11-19
PRIOR PLING DATE: 2001-11-19
PRIOR PLING DATE: 2001-10-16
PRIOR PLING DATE: 2001-06-15
PRIOR PLING DATE: 2001-06-15
PRIOR PLING DATE: 2001-06-15
PRIOR PLING DATE: 2001-06-15
PRIOR PLING DATE: 2001-03-21
PRIOR PLING DATE: 2001-03-21
PRIOR PLING DATE: 2001-03-16
PRIOR PLING DATE: 2001-03
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TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
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Pred. No. 5.3e-40;
4; Mismatches 14; Indels
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Best Local Similarity 84.4%;
Matches 108; Conservative 4
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COGANISM: Homo sapiens
US-11-054-515-981
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ORGANISM: Homo sapiens
PEATURE:
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LENGTH: 125
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US-11-054-515-1102
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                                                                                                                                                                   1 EVQLLESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGRGLEWVALIWYDGGNKEY 60
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Publication No. US2005025532A1
SEGNERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REPERBNCE: PF523P3
CURRENT PAPLICATION NUMBER: US/11/054,515
CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATEMENT INFORMATION:

APPLICANT: Kellermann, Sirid-Ai
APPLICANT: Green, Larry L.

APPLICANT: Green, Larry L.

APPLICANT: Green, Larry L.

TITLE OF INVENTION: REDUCING THE RISK OF HUMAN ANTI-HUMAN
TITLE OF INVENTION: ANTIBODIES THROUGH V GENE MANIPULATION
FILE REPERRORE ABGRAXI.100A
CURRENT APPLICATION NUMBER: US/11/084,554
CURRENT PILING DATE: 2005-03-17
PRIOR PPLING DATE: 2004-03-19
PRIOR FILING DATE: 2004-03-19
PRIOR FILING DATE: 2004-05-24
NUMBER OF SEQ ID NOS: 266

SOFTWARE: PARESEQ FOR WINDOWS VERSION 4.0

SEQ ID NOS: 266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 7; Length 125;
                                                           Query Match 82.1%; Score 548.5; DB 7; Length 125; Best Local Similarity 83.3%; Pred. No. 3e-40; Matches 105; Conservative 9; Mismatches 11; Indels 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            82.0%; Score 547.5; DB 7; Length 1
86.5%; Pred. No. 3.6e-40;
iive 3; Mismatches 13; Indels
; OTHER INFORMATION: anti-Rh(D) antibody clone SH16
US-11-064-174-140
                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 8, Application US/11084554 Publication No. US20050260679A1 GENERAL INFORMATION:
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
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US-11-054-515-397
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61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGNYYYYGMDVWGQGT 120
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PRIOR FILING DATE: 2004-02-11
PRIOR PLILING DATE: 2004-02-11
PRIOR PLILING DATE: 2004-06-18
PRIOR PLILING DATE: 2002-11-14
PRIOR PLILING DATE: 2002-11-14
PRIOR PLILING DATE: 2001-11-16
PRIOR PLILING DATE: 2001-11-16
PRIOR PLILING DATE: 2001-12-19
PRIOR PLILING DATE: 2001-12-19
PRIOR PLILING DATE: 2001-06-15
PRIOR PLILING DATE: 2001-06-15
PRIOR PLILING DATE: 2001-05-25
PRIOR PLILING DATE: 2001-05-25
PRIOR PLILING DATE: 2001-03-21
PRIOR PLILING DATE: 2001-03-21
PRIOR PLILING DATE: 2001-03-16
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TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523P3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 249;
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Pred. No. 9.2e-40;
3; Mismatches 17;
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CURRENT PELICATION NUMBER: US/11/054,515
CURRENT FILING DATE: 2005-02-10
PRIOR FILING DATE: 2006-02-11
PRIOR FILING DATE: 2004-02-11
PRIOR FILING DATE: 2004-06-18
PRIOR FILING DATE: 2004-06-18
PRIOR FILING DATE: 2004-06-18
PRIOR FILING DATE: 2001-10-14
PRIOR FILING DATE: 2001-11-16
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PRIOR FILING DATE: 2001-12-19
PRIOR FILING DATE: 2001-12-19
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PRIOR PRILING DATE: 2001-03-16
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Best Local Similarity 84.1
Matches 106; Conservative
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US-11-054-515-397
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TITLE OP INVENTION: Antibodies that Immunospecifically Bind BLyS

TITLE OP INVENTION: Antibodies that Immunospecifically Bind BLyS

FILE REPERENCE: PF523P3

CURRENT APPLICATION NUMBER: US/11/054,515

CURRENT PILING DATE: 2005-02-10

PRIOR PILING DATE: 2004-06-18

PRIOR PILING DATE: 2004-06-18

PRIOR PILING DATE: 2002-11-14

PRIOR PILING DATE: 2002-11-14

PRIOR PILING DATE: 2001-11-16

PRIOR PILING DATE: 2001-11-16

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PRIOR PILING DATE: 2001-06-15

PRIOR PILING DATE: 2001-03-21

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PRIOR PILING DATE: 2001-03-16

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                       Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 1102
LENGTH: 249
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Pred. No. 9.2e-40;
3; Mismatches 17; Indels
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Best Local Similarity 84.1%; Pred. No. 9.2e-40;
Matches 106; Conservative 3; Mismatches 17,
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Best Local Similarity 84.1%;
Matches 106; Conservative
PRIOR FILING DATE: 2000-10-17
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; ORGANISM: Homo sapiens
US-11-054-515-1115
                                                                                                                                         TYPE: PRT
ORGANISM: Homo sapiens
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61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGNYYYYGMDVWGQGT 120

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                                                                                      Sequence 1724, Application US/11054515; Publication No. US20050255532A1; GENERAL INFORMATION:
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US-11-054-515-1724
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STIC-Biotech/ChemLib

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Seharaseyon, Jegatheesan

Sent:

Tuesday, December 27, 2005 3:25 PM

To:

STIC-Biotech/ChemLib

Subject:

Re:09/920137

Importance:

High

Hi,

Please search SEQ ID NO: 7 and 8 of 09/920137 in both the commercial and interference databases.

Thanks in advance, Seyon.

J.Seharaseyon Art Unit 1647 Remsen 4C61 Mailbox 4C70

Phone: (571)-272-0892 Fax: (571)-273-0892 ME

Searcher:	
Searcher Phone:	
Date Searcher Picked up:_	
Date completed:	
Searcher Prep Time:	
Online Time:	

Type of Search		
NA#	AA#:	
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Vendors and cost where applicable STN:

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